

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: March 30, 2004, 15:07:28 ; Search time 0.513208 Seconds
(without alignments)
1609.516 Million cell updates/sec

Title: US-09-893-371A-2

Perfect score: 78

Sequence: 1 MYRASALGSDGVRVT 16

Scoring table: BLOSUM62

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Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0

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Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents AA:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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7	49	62.8	270	4	US-09-489-039A-7939
8	48	61.5	275	4	US-09-543-681A-4697
9	45	57.7	1059	4	US-09-489-039A-10044
10	44	56.4	226	4	US-09-438-144-1
11	44	56.4	268	4	US-09-438-144-7
12	44	56.4	272	4	US-09-252-991A-30277
13	41	52.6	330	4	US-09-252-991A-25521
14	40	51.3	492	4	US-09-489-039A-9018
15	39	50.0	224	4	US-09-424-978B-2
16	38	48.7	223	4	US-09-543-681A-6559
17	38	48.7	308	3	US-09-347-803-12
18	38	48.7	411	4	US-09-252-991A-21815
19	38	48.7	985	5	PCT-US96-03916-6
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21	37	47.4	270	4	US-09-252-991A-24974
22	37	47.4	298	4	US-09-252-991A-22627
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24	37	47.4	364	4	US-09-252-991A-27209
25	37	47.4	623	3	US-09-041-991A-6
26	37	47.4	623	4	US-09-608-533A-6
27	37	47.4	625	4	US-09-661-322A-48

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29	37	47.4	943	4	US-09-397-885-5
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31	36	46.2	96	4	US-09-673-395A-418
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33	36	46.2	219	4	US-09-438-144-4
34	36	46.2	251	4	US-09-438-144-6
35	36	46.2	259	4	US-09-498-520A-52
36	36	46.2	260	4	US-09-134-001C-4009
37	36	46.2	265	4	US-09-489-039A-11121
38	36	46.2	301	4	US-08-311-731A-156
39	36	46.2	361	4	US-08-173-508-2
40	36	46.2	537	2	US-08-265-310-2
41	36	46.2	537	3	US-08-951-742-2
42	36	46.2	632	4	US-09-661-322A-2
43	36	46.2	632	4	US-09-041-991A-4
44	36	46.2	635	3	US-09-608-533A-4
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ALIGNMENTS

RESULT 1

US-09-328-352-8211
; Sequence 8211, Application US/09328352
; Patent No. 6562958

; GENERAL INFORMATION:

; APPLICANT: Gary L. Breton et al.

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; FILE REFERENCE: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: GTC99-03PA

; CURRENT APPLICATION NUMBER: US/09/328,352

; CURRENT FILING DATE: 1999-06-04

; NUMBER OF SEQ ID NOS: 8252

; SEQ ID NO 8211

; LENGTH: 303

; TYPE: PRT

; ORGANISM: Acinetobacter baumannii

US-09-328-352-8211

Query Match 66.7%; Score 52; DB 4; Length 303;

Best Local Similarity 66.7%; Pred. No. 0.28;

Matches 10; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

Qy 1 MYRASALGSDGVRV 15

Db 209 VRYLASSLGVDGIRV 223

Sequence 8211, Ap

Sequence 2420, Ap

Sequence 3, Appli

Sequence 2, Appli

Sequence 8, Appli

Sequence 9, Appli

Sequence 7939, Ap

Sequence 4697, Ap

Sequence 10044, A

Sequence 1, Appli

Sequence 7, Appli

Sequence 30277, A

Sequence 25521, A

Sequence 9018, Ap

Sequence 2, Appli

Sequence 6359, Ap

Sequence 12, Appli

Sequence 21815, A

Sequence 6, Appli

Sequence 65, Appli

Sequence 24974, A

Sequence 22627, A

Sequence 3039, Ap

Sequence 27209, A

Sequence 6, Appli

Sequence 48, Appli

US-09-540-236-2420
; Sequence 2420, Application US/09540236
; Patent No. 6673910

; GENERAL INFORMATION:

; APPLICANT: Gary L. Breton et al.

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO MORAXELLA CAT

; FILE REFERENCE: 2709.2005-001

; CURRENT APPLICATION NUMBER: US/09/540,236

; CURRENT FILING DATE: 2000-04-04

; NUMBER OF SEQ ID NOS: 3840

; SEQ ID NO 2420

; LENGTH: 301

; TYPE: PRT

; ORGANISM: M.cattarrhalis

US-09-540-236-2420

Query Match 64.1%; Score 50; DB 4; Length 301;

Best Local Similarity 60.0%; Pred. No. 0.61;

Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

Qy 1 MYRASALGSDGVRV 15

RESULT 5
US-08-241-766-8
Sequence 8, Application US/08241766
Patent No. 5686590
GENERAL INFORMATION:
APPLICANT: JACOBS, W. R.
APPLICANT: COLLINS, D. M.

RESULT 5
 US-08-241-766-8
 Sequence 8, Application US/08241766
 Patent No. 5686590
 GENERAL INFORMATION:
 APPLICANT: JACOBS, W. R.
 APPLICANT: COLLINS, D. M.

GENERAL INFORMATION:
APPLICANT: JACOBS, W. R.
APPLICANT: COLLINS, D. M.
APPLICANT: BANERJEE, A.
APPLICANT: DELISLE, G. W.
APPLICANT: WILSON, T. M.
TITLE OF INVENTION: METHODS AND C
TITLE OF INVENTION: AND TREATING
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0
CURRENT APPLICATION DATA:

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Query Match      62.8%; Score 49; DB 1; Length 262;
Best Local Similarity 60.0%; Pred.No. 0.78;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY      1 MRYRASALGSDGVRV 15
      :|||:|:|:|:|:|
DB      170 VRYWANAMGPEGVRV 184
      :|||:|:|:|:|:|

RESULT 6
US-08-241-766e-9
; Sequence 9, Application US/08241766
; Patent No. 5686590
; GENERAL INFORMATION:
; APPLICANT: JACOBS, W. R.
; APPLICANT: COLLINS, D. M.
; APPLICANT: BANERJEE, A.
; APPLICANT: delISLE, G. W.
; APPLICANT: WILSON, T. M.
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DETECTING
; TITLE OF INVENTION: AND TREATING MYCOBACTERIAL INFECTIONS USING AN inha AGENT
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 Page Mill Road
; CITY: Palo Alto
; STATE: CA
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:

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;; APPLICATION NUMBER: US/08/241,766
;; FILING DATE: 12-MAY-1994
;; CLASSIFICATION: 514
;; ATTORNEY/AGENT INFORMATION:
;; NAME: MONROY, GLADYS H.
;; REGISTRATION NUMBER: 32,430
;; REFERENCE/DOCKET NUMBER: 25237-20003.20
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (415) 813-5600
;; TELEFAX: (415) 494-0792
;; TELEX: 706141
;; INFORMATION FOR SEQ ID NO: 9:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 262 amino acids
;; TYPE: amino acid
;; STRANDEDNESS: single
;; TOPOLOGY: linear
US-08-241-766-9

Query Match 62.8%; Score 49; DB 1; Length 262;
Best Local Similarity 60.0%; Pred. No. 0.78;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 15
Db 170 VRYMANAMGPEGVRV 184

RESULT 7
US-09-489-039A-7939
; Sequence 7939, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
; APPLICANT: Gary Breton et. al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
; FILE REFERENCE: 2709.2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 7939
; LENGTH: 270
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-7939

Query Match 62.8%; Score 49; DB 4; Length 270;
Best Local Similarity 60.0%; Pred. No. 0.81;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 15
Db 178 VRYMANAMGPEGVRV 192

RESULT 8
US-09-543-681A-4697
; Sequence 4697, Application US/09543681A
; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; FILE REFERENCE: 2709.1002-001
; CURRENT APPLICATION NUMBER: US/09/543,681A
; CURRENT FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
; SEQ ID NO 4697
; LENGTH: 275

;; TYPE: PRT
;; ORGANISM: Proteus mirabilis
US-09-543-681A-4697

Query Match 61.5%; Score 48; DB 4; Length 275;
Best Local Similarity 53.3%; Pred. No. 1.2;
Matches 8; Conservative 5; Mismatches 2; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 15
Db 183 VRYMANAMGPEGVRV 197

RESULT 9
US-09-489-039A-10044
; Sequence 10044, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
; APPLICANT: Gary Breton et. al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
; FILE REFERENCE: 2709.2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 10044
; LENGTH: 1059
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-10044

Query Match 57.7%; Score 45; DB 4; Length 1059;
Best Local Similarity 62.5%; Pred. No. 18;
Matches 10; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 16
Db 86 MYMKSVAGSDGLVT 101

RESULT 10
US-09-438-144-1
; Sequence 1, Application US/09438144
; Patent No. 6531291
; GENERAL INFORMATION:
; APPLICANT: Kabbash, Christina
; APPLICANT: Silverstein, Samuel C.
; APPLICANT: Shuman, Howard A.
; APPLICANT: Blanchard, Josh S.
; TITLE OF INVENTION: NOVEL ANTIMICROBIAL ACTIVITY OF GENFIBROZIL AND RELATED COMPOUNDS
; FILE REFERENCE: 0575/58043
; CURRENT APPLICATION NUMBER: US/09/438,144
; CURRENT FILING DATE: 1999-11-10
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: Patent in version 3.1
; SEQ ID NO 1
; LENGTH: 226
; TYPE: PRT
; ORGANISM: L. pneumophila
US-09-438-144-1

Query Match 56.4%; Score 44; DB 4; Length 226;
Best Local Similarity 53.3%; Pred. No. 4.8;
Matches 8; Conservative 5; Mismatches 2; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 15
Db 177 VRYLAASLGSRGURI 191

RESULT 11

US-09-438-144-7
; Sequence 7, Application US/09438144
; Patent No. 6531291
; GENERAL INFORMATION:
; APPLICANT: Kabbash, Christina
; APPLICANT: Silverstein, Samuel C.
; APPLICANT: Shuman, Howard A.
; APPLICANT: Blanchard, Josh S.
; TITLE OF INVENTION: NOVEL ANTIMICROBIAL ACTIVITY OF GEMFIBROZIL AND RELATED COMPOUNDS
; TITLE OF INVENTION: DERIVATIVES AND METABOLITES THEREOF
; FILE REFERENCE: 0575/58043
; CURRENT APPLICATION NUMBER: US/09/438,144
; CURRENT FILING DATE: 1999-11-10
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 268
; TYPE: PRT
; ORGANISM: L. pneumophila enoyl reductase
US-09-438-144-7

Query Match 56.4%; Score 44; DB 4; Length 268;
Best Local Similarity 53.3%; Pred. No. 5.8;
Matches 8; Conservative 5; Mismatches 2; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 15
Db 177 VRYLAASLGSRLRI 191

RESULT 12

US-09-252-991A-30277
; Sequence 30277, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 30277
; LENGTH: 272
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-30277

Query Match 56.4%; Score 44; DB 4; Length 272;
Best Local Similarity 53.3%; Pred. No. 5.9;
Matches 8; Conservative 4; Mismatches 3; Indels 0; Gaps 0;

QY 1 MYRASALGSDGVRV 15
Db 180 VRYLAGSLGAEGTRV 194

RESULT 13

US-09-252-991A-25521
; Sequence 25521, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18

; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 25521
; LENGTH: 330
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-25521

Query Match 52.6%; Score 41; DB 4; Length 330;
Best Local Similarity 58.3%; Pred. No. 24;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 2 RYRASALGSDGV 13
Db 316 KYRANAAGDGV 327

RESULT 14

US-09-489-039A-9018
; Sequence 9018, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
; APPLICANT: Gary Breton et. al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
; TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 2709.2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 9018
; LENGTH: 492
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-9018

Query Match 51.3%; Score 40; DB 4; Length 492;
Best Local Similarity 53.8%; Pred. No. 55;
Matches 7; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

QY 1 MYRASALGSDGV 13
Db 365 MRYPVTMGSDGL 377

RESULT 15

US-09-424-978B-2
; Sequence 2, Application US/09424978B
; Patent No. 6684445
; GENERAL INFORMATION:
; APPLICANT: Falco, Saverio Carl
; APPLICANT: Allen, Stephen M.
; APPLICANT: Rafalski, J. Antoni
; APPLICANT: Hitz, William D.
; APPLICANT: Kinney, Anthony J.
; APPLICANT: Abell, Lynne N.
; APPLICANT: Thorpe, Catherine J.
; TITLE OF INVENTION: Plant Amino Acid Biosynthetic Enzymes
; FILE REFERENCE: BB-1087
; CURRENT APPLICATION NUMBER: US/09/424,978B
; CURRENT FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: US 60/048,771
; PRIOR FILING DATE: 1997-06-06
; NUMBER OF SEQ ID NOS: 43
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2
; LENGTH: 224
; TYPE: PRT
; ORGANISM: Zea mays

Query Match 51.3%; Score 40; DB 4; Length 492;
Best Local Similarity 53.8%; Pred. No. 55;
Matches 7; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

QY 1 MYRASALGSDGV 13
Db 365 MRYPVTMGSDGL 377

US-09-424-978B-2

Query Match 50.0%; Score 39; DB 4; Length 224;
Best Local Similarity 66.7%; Pred. No. 35;
Matches 8; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

QY 4 RASALGSDGVRV 15
| | | | |
Db 153 RGQLLGEDGVRV 164

Search completed: March 30, 2004, 15:13:37
Job time : 1.51321 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: March 30, 2004, 15:11:54 ; Search time 1.19245 Seconds
(without alignments)
3510.905 Million cell updates/sec

Title: US-09-893-371A-2

Perfect score: 78

Sequence: 1 MRYRASALGSDGVRVT 16

Scoring table: BLOSUM62

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Total number of hits satisfying chosen parameters: 1065169

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Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	78	100.0	16	14	US-10-287-216-2
2	78	100.0	1044	14	US-10-287-216-1
3	56	71.8	20	14	US-10-287-216-4
4	52	66.7	288	12	US-10-282-122A-45237
5	51	65.4	262	15	US-10-369-493-388
6	51	65.4	262	15	US-10-369-493-21191
7	50	64.1	260	12	US-10-282-122A-66916
8	50	64.1	274	12	US-10-282-122A-62934
9	49	62.8	219	14	US-10-366-686-2
10	49	62.8	219	14	US-10-366-686-3
11	49	62.8	262	9	US-09-815-242-10152
12	49	62.8	262	12	US-10-282-122A-42969
13	49	62.8	262	12	US-10-282-122A-55624
14	49	62.8	262	12	US-10-282-122A-53996
15	49	62.8	262	12	US-10-282-122A-74993

16	49	62.8	262	12	US-10-282-122A-76000	Sequence 76000, A
17	49	62.8	262	15	US-10-369-493-812	Sequence 812, App
18	49	62.8	262	16	US-10-377-250-3	Sequence 3, Appli
19	49	62.8	269	9	US-09-815-242-13824	Sequence 13824, A
20	48	61.5	262	12	US-10-282-122A-68608	Sequence 68608, A
21	46	59.0	264	15	US-10-369-493-13790	Sequence 13790, A
22	45	57.7	256	15	US-10-369-493-19566	Sequence 19566, A
23	45	57.7	784	12	US-10-282-122A-55466	Sequence 55466, A
24	45	57.7	1050	12	US-10-282-122A-60043	Sequence 60043, A
25	44	56.4	190	12	US-10-424-599-202305	Sequence 202305,
26	44	56.4	227	14	US-10-366-686-1	Sequence 1, Appli
27	44	56.4	264	15	US-10-369-493-18839	Sequence 18839, A
28	44	56.4	265	9	US-09-815-242-11851	Sequence 11851, A
29	44	56.4	265	12	US-10-282-122A-66330	Sequence 66330, A
30	44	56.4	268	12	US-10-282-122A-61559	Sequence 61559, A
31	44	56.4	268	14	US-10-366-686-7	Sequence 7, Appli
32	43	55.1	252	15	US-10-369-493-4271	Sequence 4271, Ap
33	43	55.1	259	15	US-10-369-493-21833	Sequence 21833, A
34	43	55.1	263	12	US-10-282-122A-49065	Sequence 49065, A
35	43	55.1	263	12	US-10-282-122A-50662	Sequence 50662, A
36	43	55.1	323	12	US-10-425-114-64219	Sequence 64219, A
37	42	53.8	264	12	US-10-282-122A-69729	Sequence 69729, A
38	42	53.8	264	15	US-10-369-493-7881	Sequence 7881, Ap
39	41	52.6	256	15	US-10-369-493-8921	Sequence 8921, Ap
40	41	52.6	256	15	US-10-369-493-10656	Sequence 10656, A
41	41	52.6	261	12	US-10-282-122A-51117	Sequence 51117, A
42	41	52.6	451	14	US-10-128-714-3277	Sequence 3277, Ap
43	41	52.6	512	14	US-10-156-761-9793	Sequence 9793, Ap
44	41	52.6	559	14	US-10-128-714-8277	Sequence 8277, Ap
45	40	51.3	231	12	US-10-282-122A-47731	Sequence 47731, A

ALIGNMENTS

RESULT 1
US-10-287-216-2
; Sequence 2, Application US/10287216
; Publication No. US20030186375A1
; GENERAL INFORMATION:
; APPLICANT: de Lanerolle, Primal
; APPLICANT: No. US20030186375A1ak, Grzegorz
; APPLICANT: Pestig-Dragovich, Lidija
; APPLICANT: Stojiljkovic, Ljuba
; APPLICANT: Hozak, Pavel
; TITLE OF INVENTION: Nuclear Myosin I B with A 16 Amino Acid N-Terminal
; TITLE OF INVENTION: Extension
; FILE REFERENCE: 30151/92399
; CURRENT APPLICATION NUMBER: US/10/287,216
; CURRENT FILING DATE: 2002-11-04
; PRIOR APPLICATION NUMBER: US/09/893,371
; PRIOR FILING DATE: 2001-06-27
; PRIOR APPLICATION NUMBER: 60/214,944
; PRIOR FILING DATE: 2000-06-29
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 16
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:N-terminal 16
; OTHER INFORMATION: amino acid extension
US-10-287-216-2

Query Match 100.0%; Score 78; DB 14; Length 16;
Best Local Similarity 100.0%; Pred. No. 5e-07;
Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MRYRASALGSDGVRVT 16
|||
Db 1 MRYRASALGSDGVRVT 16

RESULT 2
US-10-287-216-1
; Sequence 1, Application US/10287216
; Publication No. US20030186375A1
; GENERAL INFORMATION:
; APPLICANT: de Lanerolle, Primal
; APPLICANT: No. US20030186375A1ak, Grzegorz
; APPLICANT: Pestic-dragovich, Lidija
; APPLICANT: Stojiljkovic, Ljuba
; APPLICANT: Hozak, Pavel
; TITLE OF INVENTION: Nuclear Myosin I B with A 16 Amino Acid N-Terminal
; TITLE OF INVENTION: Extension
; FILE REFERENCE: 30151/92399
; CURRENT APPLICATION NUMBER: US/10/287,216
; CURRENT FILING DATE: 2002-11-04
; PRIOR APPLICATION NUMBER: US/09/893,371
; PRIOR FILING DATE: 2001-06-27
; PRIOR APPLICATION NUMBER: 60/214,944
; PRIOR FILING DATE: 2000-06-29
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 1044
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Nuclear Myosin
US-10-287-216-1

Query Match 100.0%; Score 78; DB 14; Length 1044;
Best Local Similarity 100.0%; Pred. No. 5.2e-05;
Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MRYASALGSDGVRVT 16
DB 1 MRYASALGSDGVRVT 16

RESULT 3
US-10-287-216-4
; Sequence 4, Application US/10287216
; Publication No. US20030186375A1
; GENERAL INFORMATION:
; APPLICANT: de Lanerolle, Primal
; APPLICANT: No. US20030186375A1ak, Grzegorz
; APPLICANT: Pestic-dragovich, Lidija
; APPLICANT: Stojiljkovic, Ljuba
; APPLICANT: Hozak, Pavel
; TITLE OF INVENTION: Nuclear Myosin I B with A 16 Amino Acid N-Terminal
; TITLE OF INVENTION: Extension
; FILE REFERENCE: 30151/92399
; CURRENT APPLICATION NUMBER: US/10/287,216
; CURRENT FILING DATE: 2002-11-04
; PRIOR APPLICATION NUMBER: US/09/893,371
; PRIOR FILING DATE: 2001-06-27
; PRIOR APPLICATION NUMBER: 60/214,944
; PRIOR FILING DATE: 2000-06-29
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 20
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:NMI Beta
; OTHER INFORMATION: Peptide Overlapping Consensus Start Site
US-10-287-216-4

Query Match 71.8%; Score 56; DB 14; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.0053;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 5 ASALGSDGVRVT 16
DB 1 ASALGSDGVRVT 12

RESULT 4
US-10-282-122A-45237
; Sequence 45237, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Chlsen, Kari
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 45237
; LENGTH: 288
; TYPE: PRT
; ORGANISM: Acinetobacter baumannii
US-10-282-122A-45237

Query Match 66.7%; Score 52; DB 12; Length 288;
Best Local Similarity 66.7%; Pred. No. 0.53;
Matches 10; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 MRYASALGSDGVRV 15
DB 194 VRYLASSLGVDGIRV 208

RESULT 5
US-10-369-493-388
; Sequence 388, Application US/10369493
; Publication No. US20030233675A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.

```

; APPLICANT: Goldman, Barry S.
; APPLICANT: Chen, Xianfeng
; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
; CURRENT FILING DATE: 2003-02-28
; PRIOR APPLICATION NUMBER: US 60/360,039
; PRIOR FILING DATE: 2002-02-21
; NUMBER OF SEQ ID NOS: 47374
; SEQ ID NO 388
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Xenorhabdus nematophilus
; NAME/KEY: unsure
; LOCATION: (1)..(262)
; OTHER INFORMATION: unsure at all Xaa locations
US-10-369-493-388

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Query Match      65.4%; Score 51; DB 15; Length 262;
Best Local Similarity 60.0%; Pred. No. 0.71;
Matches 9; Conservative 5; Mismatches 1; Indels 0; Gaps 0;

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Qy 1 MRYRASALGSDGVRV 15
Db 170 VRYMANAMGAEGVRV 184

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RESULT 6

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US-10-369-493-21191
; Sequence 21191, Application US/10369493
; Publication No. US20030233675A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Goldman, Barry S.
; APPLICANT: Chen, Xianfeng
; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
; CURRENT FILING DATE: 2003-02-28
; PRIOR APPLICATION NUMBER: US 60/360,039
; PRIOR FILING DATE: 2002-02-21
; NUMBER OF SEQ ID NOS: 47374
; SEQ ID NO 21191
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Xenorhabdus nematophilus
US-10-369-493-21191

```

```

Query Match      65.4%; Score 51; DB 15; Length 262;
Best Local Similarity 60.0%; Pred. No. 0.71;
Matches 9; Conservative 5; Mismatches 1; Indels 0; Gaps 0;

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Qy 1 MRYRASALGSDGVRV 15
Db 170 VRYMANAMGAEGVRV 184

```

RESULT 7

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US-10-282-122A-66916
; Sequence 66916, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari
; APPLICANT: Zyskind, Judith

```

```

; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Cairr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 66916
; LENGTH: 260
; TYPE: PRT
; ORGANISM: Pasteurella multocida
US-10-282-122A-66916

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Query Match      64.1%; Score 50; DB 12; Length 260;
Best Local Similarity 64.3%; Pred. No. 1.1;
Matches 9; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

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Qy 2 RYRASALGSDGVRV 15
Db 171 RFWAALGKDGIRV 184

```

RESULT 8

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US-10-282-122A-62934
; Sequence 62934, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.

```

```

; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23

```



```

; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 62934
; LENGTH: 274
; TYPE: PRT
; ORGANISM: Moraxella catarrhalis
US-10-282-122A-62934

Query Match      64.1%; Score 50; DB 12; Length 274;
Best Local Similarity 60.0%; Pred. No. 1.1;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 1 MRYRASALGSDGVRV 15
Db 175 VRYLASSLGEGIRV 189

RESULT 9
US-10-366-686-2
; Sequence 2, Application US/10366686
; Publication No. US20030191146A1
; GENERAL INFORMATION:
; APPLICANT: Kabbash, Christina
; APPLICANT: Silverstein, Samuel
; APPLICANT: Shuman, Howard A
; APPLICANT: Blanchard, John S
; TITLE OF INVENTION: NOVEL ANTIMICROBIAL ACTIVITY OF GEMFIBROZIL AND RELATED COMPOUNDS
; FILE REFERENCE: 0575/58043
; CURRENT APPLICATION NUMBER: US/10/366,686
; CURRENT FILING DATE: 2003-02-13
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 2
; LENGTH: 219
; TYPE: PRT
; ORGANISM: E. coli
US-10-366-686-2

Query Match      62.8%; Score 49; DB 14; Length 219;
Best Local Similarity 60.0%; Pred. No. 1.3;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 1 MRYRASALGSDGVRV 15
Db 170 VRYMANAMGPEGVRV 184

RESULT 10
US-10-366-686-3
; Sequence 3, Application US/10366686
; Publication No. US20030191146A1
; GENERAL INFORMATION:
; APPLICANT: Kabbash, Christina
; APPLICANT: Silverstein, Samuel
; APPLICANT: Shuman, Howard A

```

```

; APPLICANT: Blanchard, John S
; TITLE OF INVENTION: NOVEL ANTIMICROBIAL ACTIVITY OF GEMFIBROZIL AND RELATED COMPOUNDS
; FILE REFERENCE: 0575/58043
; CURRENT APPLICATION NUMBER: US/10/366,686
; CURRENT FILING DATE: 2003-02-13
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 3
; LENGTH: 219
; TYPE: PRT
; ORGANISM: S. typhimurium
US-10-366-686-3

Query Match      62.8%; Score 49; DB 14; Length 219;
Best Local Similarity 60.0%; Pred. No. 1.3;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 1 MRYRASALGSDGVRV 15
Db 170 VRYMANAMGPEGVRV 184

RESULT 11
US-09-815-242-10152
; Sequence 10152, Application US/09815242
; Patent No. US20020061569A1
; GENERAL INFORMATION:
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari L.
; APPLICANT: Zyskind, Judith W.
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John D.
; APPLICANT: Carr, Grant J.
; APPLICANT: Yamamoto, Robert T.
; APPLICANT: Xu, H. Howard
; TITLE OF INVENTION: Identification of Essential Genes in
; FILE REFERENCE: ELITRA.011a
; CURRENT APPLICATION NUMBER: US/09/815,242
; CURRENT FILING DATE: 2001-03-21
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; NUMBER OF SEQ ID NOS: 14110
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10152
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Escherichia coli
US-09-815-242-10152

Query Match      62.8%; Score 49; DB 9; Length 262;
Best Local Similarity 60.0%; Pred. No. 1.6;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 1 MRYRASALGSDGVRV 15
Db 170 VRYMANAMGPEGVRV 184

RESULT 12

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US-10-282-122A-42969
; Sequence 42969, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 42969
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Escherichia coli
US-10-282-122A-42969

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Query Match 62.8%; Score 49; DB 12; Length 262;
Best Local Similarity 60.0%; Pred. No. 1.6;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

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```

Qy 1 MRYRASALGSDGVRV 15
:|||||:|:|:|
Db 170 VRYMANAMGPGVRV 184

```

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RESULT 13
US-10-282-122A-55624
; Sequence 55624, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert

```

```

; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 55624
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Enterobacter cloacae
US-10-282-122A-55624

```

```

Query Match 62.8%; Score 49; DB 12; Length 262;
Best Local Similarity 60.0%; Pred. No. 1.6;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

```

```

Qy 1 MRYRASALGSDGVRV 15
:|||||:|:|:|
Db 170 VRYMANAMGPGVRV 184

```

```

RESULT 14
US-10-282-122A-59396
; Sequence 59396, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06

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; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 59396
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-10-282-122A-59396

Query Match          62.8%; Score 49; DB 12; Length 262;
Best Local Similarity 60.0%; Pred. No. 1.6;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

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Db      170 VRYMANMGPEGVRV 184

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; Sequence 74993, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
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; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 74993
; LENGTH: 262
; TYPE: PRT
; ORGANISM: Salmorella typhimurium
US-10-282-122A-74993

Query Match          62.8%; Score 49; DB 12; Length 262;
Best Local Similarity 60.0%; Pred. No. 1.6;
Matches 9; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

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Db      170 VRYMANMGPEGVRV 184

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: March 30, 2004, 15:11:54 ; Search time 77.8075 Seconds
(without alignments)
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Title: US-09-893-371A-1
Perfect score: 5398
Sequence: 1 MRYRASALGSDGVRVTWESA.....LITKAKNGLAVVAPRLNSR 1044

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1065169 seqs, 261661801 residues

Total number of hits satisfying chosen parameters: 1065169

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA:
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18: /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	5398	100.0	1044	14	US-10-287-216-1
2	2119	39.3	1078	15	Sequence 1, Appli
3	1756	32.5	1017	15	Sequence 24, Appl
4	1745	32.3	1013	12	Sequence 5684, Ap
5	1745	32.3	1013	12	US-10-369-493-5684
6	1745	32.3	1013	12	US-10-092-900A-230
7	1732	32.1	688	15	Sequence 128, App
8	1730.5	32.1	1018	12	Sequence 128, App
9	1730.5	32.1	1018	12	Sequence 152, App
10	1722.5	31.9	1098	15	Sequence 164, App
11	1714.5	31.8	1098	15	Sequence 126, App
12	1709.5	31.7	1096	12	Sequence 154, App
13	1709.5	31.7	1096	12	US-10-040-047-3280
14	1594.5	29.5	670	12	Sequence 2, Appli
15	1564	29.0	1100	15	Sequence 26, Appl
					Sequence 66, Appl
					Sequence 165, App
					Sequence 5164, Ap

16	1530.5	28.4	734	15	US-10-028-248A-108
17	1530.5	28.4	734	15	US-10-107-782-108
18	1509.5	28.0	1216	15	US-10-369-493-2390
19	1503	27.8	1258	15	US-10-369-493-3891
20	1501	27.8	1219	15	US-10-369-493-1885
21	1450	26.9	1271	15	US-10-369-493-22237
22	1392.5	25.8	1234	15	US-10-369-493-13287
23	1357	25.1	2202	15	US-10-094-466-12
24	1285.5	23.8	2058	10	US-09-815-379-17
25	1285.5	23.8	2058	12	US-10-211-462-207
26	1285.5	23.8	2058	14	US-10-021-660-120
27	1270	23.5	1742	15	US-10-012-697-1548
28	1261.5	23.4	1016	13	US-10-044-303-2
29	1261	23.4	1537	15	US-10-369-493-22785
30	1241.5	23.0	2048	10	US-09-815-379-12
31	1241.5	23.0	2057	10	US-09-815-379-10
32	1240	22.5	765	13	US-10-044-303-1
33	1216	22.5	1019	15	US-10-369-493-6432
34	1216	22.5	1855	14	US-10-177-293-315
35	1199	22.2	1956	15	US-10-369-493-6729
36	1192	22.1	1938	14	US-10-171-311-164
37	1192	22.1	1972	14	US-10-171-311-162
38	1192	22.1	1972	15	US-10-341-434-103
39	1190	22.0	1474	15	US-10-369-493-2556
40	1188.5	22.0	1945	10	US-09-927-597-2
41	1188.5	22.0	1979	10	US-09-927-597-4
42	1188	22.0	1673	15	US-10-085-198-132
43	1187	22.0	1524	12	US-10-336-472-18
44	1187	22.0	1859	12	US-10-336-472-20
45	1187	22.0	1935	12	US-10-336-472-22

ALIGNMENTS

RESULT 1
US-10-287-216-1
; Sequence 1, Application US/10287216
; Publication No. US20030186375A1
; GENERAL INFORMATION:
; APPLICANT: de Lanerolle, Primal
; APPLICANT: No. US20030186375A1ak, Grzegorz
; APPLICANT: Pestic-Dragovich, Lidija
; APPLICANT: Stojiljkovic, Ljuba
; APPLICANT: Horak, Pavel
; TITLE OF INVENTION: Nuclear Myosin I B with A 16 Amino Acid N-Terminal
; TITLE OF INVENTION: Extension
; FILE REFERENCE: 30151/92399
; CURRENT APPLICATION NUMBER: US/10/287,216
; CURRENT FILING DATE: 2002-11-04
; PRIOR APPLICATION NUMBER: US/09/893,371
; PRIOR FILING DATE: 2001-06-27
; PRIOR APPLICATION NUMBER: 60/214,944
; PRIOR FILING DATE: 2000-06-29
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 1044
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Nuclear Myosin
; OTHER INFORMATION: 1 beta
US-10-287-216-1

Query Match 100.0%; Score 5398; DB 14; Length 1044;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1044; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MRYRASALGSDGVRVTWESALTARDVGQDFVLLNFSTSEAAFIENRRFRRLIYTY 60
Db 1 MRYRASALGSDGVRVTWESALTARDVGQDFVLLNFSTSEAAFIENRRFRRLIYTY 60

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QY 121 GESGAGKTATKRLLOFYAETCPAPRGGAVERDLQSNPVLFAFGNAKTLRNDSSRF 180
Db 121 GESGAGKTATKRLLOFYAETCPAPRGGAVERDLQSNPVLFAFGNAKTLRNDSSRF 180
QY 181 KYMDVDFPKGAPVGGHILSYLLEKSRVHQHNGENFHVYQLLEGGEETLRILGLER 240
Db 181 KYMDVDFPKGAPVGGHILSYLLEKSRVHQHNGENFHVYQLLEGGEETLRILGLER 240
QY 241 NFQSYLYLVKGCAKVSINDSKDWMKALSVIDFTEDEVEDLLSIVASVLHGNHF 300
Db 241 NFQSYLYLVKGCAKVSINDSKDWMKALSVIDFTEDEVEDLLSIVASVLHGNHF 300
QY 301 ADEDSNAQVTENQKYLTRLLGVEGTTLRALTHRKIIAKGEBLLSPNLREQAAYARD 360
Db 301 ADEDSNAQVTENQKYLTRLLGVEGTTLRALTHRKIIAKGEBLLSPNLREQAAYARD 360
QY 361 ALAKAVYSRTFTWLVKINRSKASDAESPWSRSTTVLGLLDIYGFEVFOHNSRPOFCIN 420
Db 361 ALAKAVYSRTFTWLVKINRSKASDAESPWSRSTTVLGLLDIYGFEVFOHNSRPOFCIN 420
QY 421 YCNEKLQQLFIELTLKSEQEEYEAEGIAWEPVQYFNKKIICDLVBEKFKGIISILDECL 480
Db 421 YCNEKLQQLFIELTLKSEQEEYEAEGIAWEPVQYFNKKIICDLVBEKFKGIISILDECL 480
QY 481 RGEATDLTFLEKLEDTVPKPHFLTHKLADQKTRKSLDRGFRLLHAGVTVYVTGFL 540
Db 481 RGEATDLTFLEKLEDTVPKPHFLTHKLADQKTRKSLDRGFRLLHAGVTVYVTGFL 540
QY 541 DXNNLLFNLKETWCSSNMPMAOCFDSKSELDKRRPETVATQKMSLLQLVEILRSKE 600
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QY 601 PAYIRCIKENDAKQPGREDEVILRHQVKYGLGLMENLVRERAGFAYRRKYEAPLQYKSLC 660
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QY 661 PETPMWAGRPQDGVAVLVRHLGYKPEEYKMGRTKIFIRPKTLFATEDSLEVRQSLAT 720
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Db 841 CRSISPENKQLOKAVASEIFKGGKDNYPQSVPLFISTRIGTBEISPRVLQSLGSEPI 900
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QY 961 HVQEDNKQKGVVLQSDHVIETLTKTALSADRVNNINQGSITTFAGPGRDGIIDFTS 1020
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RESULT 2

US-10-144-194A-24

; Sequence 24, Application US/10144194A

; Publication No. US20030215809A1

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; GENERAL INFORMATION:
; APPLICANT: Origene Technologies Inc
; TITLE OF INVENTION: Regulation: Regulated Breast Cancer Genes
; FILE REFERENCE: 3U 103 R1
; CURRENT APPLICATION NUMBER: US/10/144,194A
; CURRENT FILING DATE: 2002-06-12
; NUMBER OF SEQ ID NOS: 114
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 24
; LENGTH: 1078
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-10-144-194A-24
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Best Local Similarity 43.0%; Pred. No. 2.1e-183;

Matches 465; Conservative 184; Mismatches 343; Indels 90; Gaps 20;

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QY 87 GVSFEYVPHLFAVADTVYRALTRERDQAVMISGESGAGKTATKRLLOFYAETCPAPE 146
Db 74 NNFVELSPHIFALSDEAYRSLRDQDKQCILITGESGAGKTEASKLVMSYVAAVCGKA 133
QY 147 RGGAVRDLQSNPVLFAFGNAKTLRNDSSRFKYMVDQDFKGAPOVGGHILSYLLEKS 206
Db 134 EVNQYKEQLQSNPVLFAFGNAKTLRNDSSRFKYMIDEFDFKGDPLGGVISNYLLEKS 193
QY 207 RVHQNCHGRNFHVYQLLEGGEETLRRLGLENPQSYLYLVKGQCAKVSINDSKDWMK 266
Db 194 RVVQKQGRNFHVYQLLGSASELLNKLKLEDFSRNYL-SLDSAKVNGVDDAANFR 252
QY 267 VMKALSVIDFTEDEVEDLLSIVASVLHGNHFAADEDSN- ---AQVTENQLKYLTRL 322
Db 253 TVRNAMQIVGPMDEHAEBSVLAAVAAVLKLGNIIEFKPSRVNGLDES KIKQKELKEICEL 312
QY 323 LGVEGTTLRALTHRKIIAKGEBLLSPNLREQAAYARDALAKAVYRTFTWLVKINRS 382
Db 313 TGIDQSVLERAFSPRTVEAKQEKVSTTLNVAQAYARDALAKNLYSRFLSVLRINESI 372
QY 383 ASKDAECPWSRSTTVLGLLDIYGFEVFOHNSRFOFCINCEKLOQLFIETLKEQEEY 442
Db 373 KAQTKVRKK- ---VMGVLDIYGFEIFEDNSRFOFCINCEKLOQLFIETLKEQEEY 427
QY 443 EAGIAWEPVQYFNKKIICDLVBEKFKGIISILDEECLRPGAEATDLTFLEKLEDTVPHP 502
Db 428 IREDIEWTHIDYFNNAIICDLIENNTNGILAMLDEECLRPGVTDTETFLEKLNQVCATHQ 487
QY 503 HFLTH- ---KLADQKTRKSLDRGFEPLHLYAGEVTVYVTGFLDKNDLLFRNLKETWCSSMN 560
Db 488 HFESRMKCSRFLNDTSLPHSCFRIOHYAGKVLVQVGFVDKNDLLYRDLQSNMWKASH 547
QY 561 PMAQCFDKSELS- ---DKRPEVTATQKMSLLQLVEILRSKEPAYIRCIKENDAKQPGF 618
Db 548 ALIKSLPEGNPKNILNKRPTAGSQKASVATLMKNLQYKNFYIRCIKENDAKKAHIF 607
QY 619 DEVILRHQVKYGLGLMENLVRERAGFAYRRKYEAPLQYKSLCPETPMWAGRPQDGVAVL 678
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QY 679 VRHLGYKPEEKMGRTKIFIRPKTLFATEDSLEVRQSLATKIQAAWRGPHWQKFLVR 738
Db 668 FNELEIPEEYSFGRSKIFIRNPRTLKLEDLRQRLEDLATLIQTYRGMKCKTHFLM 727
QY 739 KRSALCIQSWWRGTLGRKKAQRKAAQTIRRLIRGFE- ---ILRHSRCPCE- --- 785
Db 728 KKSQIVLAAYRRYAQQRYQOTYKSSALVQSVIRGHWKARKILRELHQRCKRAVTTIA 787
QY 786 - - - - -NA- - - - -FFLDHVRASFLNLRRLQPLRNVLDTSWPTPPA 820
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 Db 906 QGAYL---EINKPKYKKLDAIEEKIIAEBVANKINRANGKSTSRIFLLTNNLLADQ 962
 Qy 933 -DAVKQRIDYANLTGTSVSLSDSLFVLHVQR--EDNKQKGDVVLQSDHVIETLTK--T 987
 Db 963 KSGQKSEVPVLDVTKVSMSSQNGDFFAVHULKEGSEAAKSGDFFLSSDHLIEMATKYRT 1022
 Qy 988 ALSADRVN-NINI-----NCGSITFAGGPGRDGIIIFTSGSELLIIFKAKNGHLAV 1036
 Db 1023 TLSQTKQKLNIEISDEFVLQVRQDKVCVKFQGNQKNGSV-----TCKRKNRLLLE 1074
 Qy 1037 VA 1038
 Db 1075 VA 1076

RESULT 3
 ; Sequence 5684, Application US/10369493
 ; Publication No. US20030233675A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Cao, Yongwei
 ; APPLICANT: Hinkle, Gregory J.
 ; APPLICANT: Slater, Steven C.
 ; APPLICANT: Goldman, Barry S.
 ; APPLICANT: Chen, Xianfeng
 ; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
 ; FILE REFERENCE: 38-10(52052)B
 ; CURRENT APPLICATION NUMBER: US/10/369,493
 ; CURRENT FILING DATE: 2003-02-28
 ; PRIOR APPLICATION NUMBER: US 60/360,039
 ; PRIOR FILING DATE: 2002-02-21
 ; NUMBER OF SEQ ID NOS: 47374
 ; SEQ ID NO 5684
 ; LENGTH: 1017
 ; TYPE: PRT
 ; ORGANISM: Caenorhabditis elegans
 ; US-10-369-493-5684

Query Match 32.5%; Score 1756; DB 15; Length 1017;
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Qy 28 GVQDFVLENTFSAAFTENLRFRRENLIYTYIGPVLVSNPYRDQLQYSRQHMERYRG 87
 Db 12 GVEDVLLSTTDLKSV-VQNLQRFQKGRIVTYIGEVLVAVNPYRQLGIVEKSTVDQYKG 70
 Qy 88 VSFVEVPHLPFAVDVTRALTRERRQAVMISGESGAGKTEATKRLQFVAETCPAPER 147
 Db 71 REIYERAPHVFAIAAAYRSMKRFGRDSCIIVISGAGKTETSKIIMKYLAAITNVRRQ 130
 Qy 148 G--GAVDRLLQSNPVLAEAFNAKTLRNDNSRSGKMYDVQDFKGPVGGHILSYLLEK 205
 Db 131 GEIRSVNKLRSNCILLEAFCAKTLRNDNSRSGKMYHINFDYDGPVGGNTISNYLLEK 190
 Qy 206 SRVHQHNGERNFHFVYQLLGGGEEETLRLGLERNPOSILYLVKGCACVSSINDKSDW 265
 Db 191 SRVRVQGGERNFHFVYQLVNGDDGLRQGLTKDAKQYVFLNQGGSHKVASINDSRDP 250
 Qy 266 KVMKAL-SVIDFTEDEVEDLLSVASVLHIGNTHFAADESN--AQVTENQLKYLTRL 322
 Db 251 AEVQTALRSIHTFDKQDVESMWSVIAGLIHLGNVRFDGENSGAVHIAEKAALQNAARC 310
 Qy 323 LGVEGTTLREALTHRKIIAKGEEILLSPLNLEQAAAYARDALAKAVYSFTFWLVRKINRSL 382
 Db 311 LNVTFDELAKSLSSQVAAAGDIVKKQHDVNAAYYTRDALAKALYERLFSVMVSKVNEAI 370

Qy 383 ASKDAESPSMRSTTVLGLLDIYGFEVPHQNSFQFCINCYNEKLOQLFIETLKSEOEY 442
 Db 371 SVQN--SSRYSKSHVIGVLDIYGFEIPGTNSFEOLCINCYNEKLOQLFIETLVLKQOEY 428
 Qy 443 EAEGIAWEPYQYFNKKIICDLVEEKFKGIISILDEECILRPGCEATDLTFLEKLEDTVPHP 502
 Db 429 EREGIKWVKIYFNKKVICDLVEIPRTGILSILDEACASIGNVTDKVFELGELDKLASHK 488
 Qy 503 HFLTHKLADOKTSLDRGFRLLHYAGEVTVYSGFLDKDNLLFRNLKMETKSSMNP 562
 Db 489 HYTSRNL--KQSDKSMGFEFKITHYAGDVTVYVMGFMKDKNTLFDQLRLLVHSHNRL 546
 Qy 563 MAQCF--DKSELSDKKRPETVATQFQWLSLQLVEILLRKEPAXIRCIKENDAKQPGED 619
 Db 547 VKSLFPDGSMAEVRNRPPTAGFLFNKSMSELVKQLAQKEPHYIRCIKNEEKNSVFD 606
 Qy 620 EVLIRHQVKYGLHMLNLRVRRAGFAVRRKYEAFLQRYKSLCPTETPMWAGRP-----QD 673
 Db 607 LERVEHQVRYLGLLENVRVRRAGFAHRMPYDRFVNRKYLICASTWP---NPRGQQLKD 662
 Qy 674 GVAIVRHLGYKPEEYKMGRTKIFIRPKTLFATEDSLEVRQSLATKIQAAWRGFTWRQ 733
 Db 663 SCMQILLESAGL-AQDCVQGRTKIFIRSPQTVFRLLELETEQLPNVITFLQWVRGVQORE 721
 Qy 734 KFLRVKESAICIQSWWRGTGLRRKAAKRWAAQTIRRLIRGFILRHSPRCPENAFELDHV 793
 Db 722 RY-----RMLAVRKIIGAVRRYKLSYIWO-----747
 Qy 794 RASFLMLRLRQLPRNVLDTS---WPTPPALREASSELLRELCKMNMVWYCRSIS---P 846
 Db 748 ---LINAFRDV-RMRDLGKSIRWPAPPLVLAQFVSRLRVMHQ---WRAATILARMPP 799
 Qy 847 EWKQLOQKAVASEIFKGGKDN-----YPSVPRLFISTRLCTEISPRVLQSLG 896
 Db 800 HLRASLPQKTAAPFVLNKNKNMNGYTRMWRGDLVSOQEELELPTTVSYTHDGIQALRQ-- 857
 Qy 897 SEP---IQYAVPVVKYDRKGYKPRPROLLITPSAVVIVEDAK---VKORIDYANLTGISV 950
 Db 858 SHFPGKVLFTYVQKEN-KFNKSLRVLIVTDPRFAKLENKKFKLKEPPLQGISISIV 916
 Qy 951 SSLSDSLFVLHVQRED-----NKQKGDVVLQSDHVIE 982
 Db 917 CAESNGLFVTHVGDNDIVGCAKNTKNEERVGEMIGTLAHYDKITMERSPVLIQS-AVVC 975
 Qy 983 TL---TKTALSADRVN 996
 Db 976 TLGKTKTIRVFDANN 992

RESULT 4
 US-10-092-900A-230
 ; Sequence 230, Application US/10092900A
 ; Publication No. US20040043382A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Padigaru, Muralidhara
 ; APPLICANT: Spytek, Kimberly A.
 ; APPLICANT: Shenoy, Suresh G.
 ; APPLICANT: Taupier Jr., Raymond J.
 ; APPLICANT: Pena, Carol E.A.
 ; APPLICANT: Li, Li
 ; APPLICANT: Zerhusen, Bryan D.
 ; APPLICANT: Gusev, Vladimir Y.
 ; APPLICANT: Ji, Weizhen
 ; APPLICANT: Gorman, Linda
 ; APPLICANT: Miller, Charles E.
 ; APPLICANT: Kekuda, Ramesh
 ; APPLICANT: Patturajan, Meera
 ; APPLICANT: Gangolli, Baha A.
 ; APPLICANT: Vernet, Corine A.M.
 ; APPLICANT: Guo, Xiaojia Sasha
 ; APPLICANT: Tchernev, Velizar T.
 ; APPLICANT: Fernandes, Elma R.


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; APPLICANT: Rastelli, Luca
; APPLICANT: MacDougall, John R.
; APPLICANT: Mishra, Vishnu
; APPLICANT: Pena, Carol E.A.
; APPLICANT: Spaderna, Steven K.
; APPLICANT: Shinkets, Richard A.
; APPLICANT: Smithson, Glenda
; APPLICANT: Spytek, Kimberly A.
; APPLICANT: Stone, David J.
; APPLICANT: Shenoy, Suresh G.
; APPLICANT: Ort, Tatiana
; APPLICANT: Taupier Jr, Raymond J.
; APPLICANT: Tchernev, Velizar T.
; APPLICANT: Vernet, Corine A.M.
; APPLICANT: Wolenc, Adam R.
; APPLICANT: Zerhusen, Bryan D.
; APPLICANT: Zhong, Mei
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-533C
; CURRENT APPLICATION NUMBER: US/10/336,472
; CURRENT FILING DATE: 2003-01-03
; PRIOR FILING DATE: 09/746,491
; PRIOR APPLICATION NUMBER: 10/005,041
; PRIOR FILING DATE: 2001-12-20
; PRIOR APPLICATION NUMBER: 10/023,691
; PRIOR FILING DATE: 2001-12-18
; PRIOR APPLICATION NUMBER: 10/024,212
; PRIOR FILING DATE: 2001-12-18
; PRIOR APPLICATION NUMBER: 10/055,569
; PRIOR FILING DATE: 2001-10-26
; PRIOR APPLICATION NUMBER: 10/080,334
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 10/092,900
; PRIOR FILING DATE: 2002-03-07
; PRIOR APPLICATION NUMBER: 10/136,826
; PRIOR FILING DATE: 2002-05-01
; PRIOR APPLICATION NUMBER: 10/236,417
; PRIOR FILING DATE: 2002-09-06
; PRIOR APPLICATION NUMBER: 60/345,092
; PRIOR FILING DATE: 2002-01-04
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 230
; SOFTWARE: CuraSeqList version 0.1
; SEQ ID NO 128
; LENGTH: 1013
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-10-336-472-128

Query Match 32.3%; Score 1745; DB 12; Length 1013;
Best Local Similarity 41.7%; Pred. No. 2.6e-149;
Matches 403; Conservative 155; Mismatches 339; Indels 70; Gaps 21;

QY 28 GVQDFVLENTSEAFIENLRFRFRENLIYTYIGPVLSVSNPYRDIQYSRQHMERYRG 87
Db 10 GKPDFVLLDQVTE--DFMRNLQRFEGRIYTYIGEVLSVSNPYQELPLYGPAIARYQG 68
QY 88 VSYEYVPPHLPFAVDYTRALRTERDQAVNISGESGAGKTEATKRLQFYAETCPAPER 147
Db 69 RELYERPPHLYAVANAAYKAMKHSRDTCTIVISGESGAGKTEASKHIMQYIAAVTNPSQR 128
QY 148 GGA--VRDRLLQSNPVLFAEGNAKTLENDNSRFGKYMVDQFQKAPVCGHLSVLEK 205
Db 129 AEVERVKDVLKSTCVLEAFGNARTNRNHSRFGKYMDFNDFKGPIDGGHITHSVLEK 188
QY 206 SRVYHQHGRNHFVYQLLEGGEETRLRLGLERNPQSYLYLVKGCACVSSINDKSDW 265
Db 189 SRVLKQHVGRNHFVYQLLEGGEETRLRLGLERNPQSYLYLVKGCACVSSINDKSDW 247
QY 266 KVMRKALSVIDFTEDEVEDLLSIVASVHLGNTHFAADESDNAQ-----VTTENOLKYLIT 320
Db 248 QAVTEAMRVIGFSPPEVESVHRILAILHLGNIEFVETEGGQKEGLAVAEALVDHVA 307

; APPLICANT: Agse et al.
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-442C
; CURRENT APPLICATION NUMBER: US/10/236,417
; CURRENT FILING DATE: 2003-01-06
; PRIOR APPLICATION NUMBER: US60/318,120
; PRIOR FILING DATE: 2001-09-01
; PRIOR APPLICATION NUMBER: US60/318,430
; PRIOR FILING DATE: 2001-09-10
; PRIOR APPLICATION NUMBER: US60/322,781
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/318,184
; PRIOR FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: US60/361,663
; PRIOR FILING DATE: 2002-03-05
; PRIOR APPLICATION NUMBER: US60/396,412
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US60/322,636

321 RLIGVEGTTUREALTURKIIAKGEEELSP-LINLEQAAAYARDALAKAVYSRFTTWLVRKIN 379
308 ELTATPRDLVRLSILARTVASGGRELIEKGHTAAEASAYARDACAKAVYQRLFVWVNRKIN 367
QY 380 RSLASKDAESPSRSTTVLGLLDIYGFVEFOHNSFPOPCINYNCKEKLQOLFETLKSEQ 439
Db 368 SVMPEGRDRPRDGKDTVIGVLDIYGFVEFPVNSFEQFCINYNCKEKLQOLFETLKSEQ 427
QY 440 EEEYAEAGIAWEPVQYFNKKIICDLVBEKFKGIISILDEECLRPGCEATDLTFLEKLEDTVK 499
Db 428 EEEYREGITWQSVYFNNAIVDLVERPHGILAVLDEACSSAGTITDRFLQTLDMHHR 487
QY 500 PHPEFLTHKLADQKTRKSLDRG--EFLRLHYAGVYTSVTGFLDKNDLLPRNLKBTWCSS 558
Db 488 HHLHYTSRQLC--PTDKTMEFGDRDFRIKHVAGDVTSVEGFIDKNDRDFLQDFKRLLYNS 545
QY 559 MNPIMAQCF----DKSELSDKKRPETVATQFMSLLQLVLEILRSKEPAVIRCIKPNDAK 613
Db 546 TDTILRAMPDGQDITEVT--KRPLTAGTLFKNSWVALVENLASKEPFFVRCIKPNEDK 603
QY 614 QPCRFDVLRHQVKYLGMLNLRVRRAGFAYRRKYEAFLQRYKSLCPTETWP-MWAGRPQ 672
Db 604 VAGKLDENHCHQVAVYLGLENVVRVRRAGFASRQPSRFLRYKMTCEYTPWNLHLLGSDK 663
QY 673 DGVAVLVRHLGYKPEYKMGRTKIFIRPKTLFATSDSLEVRQSLATKIQAARWGFHWR 732
Db 664 AAVSALLEHQGLQ-GDVAFGHSKLFIRSPRTLVTLEQS-----R 701
QY 733 QKELRVKRSALICISQSWRGFTLGRKAAKRWAAQTIRRLIRGFIILRHSRCPENAFELDH 792
Db 702 ARLIPI--IVLLQKAWRGTLARW--CRLRLAIYTIMRWR-----RHKVR-----HLAE 749
QY 793 VRASFLLNLRQLPRNVLDTSWPTPPPALREASELLRELCMKNMVWKYCRSISPEWKQL 852
Db 750 LQRRF--QAARQPLYGRDLVWPLPAVLQPPFDTCHALFCRWRAQLVKNIPSPDMPQI 807
QY 853 QKAVASEIFKGGKONY--POSVPRLFISTRLCTEISPRVLOS-----GSEIQA 903
Db 808 KAKVAAMGALQGLRQDMGCRRAWARDYSSATDPTASSLFAQLRKLTLQDKGFGAVLFS 867
QY 904 VPVVKYDRKGYKPRPRQLLLTFSAVVIVE--DAKYQRIDYANLGTISVSSLSDSIFVL 960
Db 868 SHVRKYNR-FHKIRNRLALLTDHLYKLDPRQYRVNRVAPLEAVTGLSVTSGDQLVVL 926
QY 961 HVQREDN 967
Db 927 HARGQDD 933

RESULT 6
US-10-236-417-152
; Sequence 152, Application US/10236417
; Publication No. US20040048256A1
; GENERAL INFORMATION:
; APPLICANT: Agse et al.
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-442C
; CURRENT APPLICATION NUMBER: US/10/236,417
; CURRENT FILING DATE: 2003-01-06
; PRIOR APPLICATION NUMBER: US60/318,120
; PRIOR FILING DATE: 2001-09-01
; PRIOR APPLICATION NUMBER: US60/318,430
; PRIOR FILING DATE: 2001-09-10
; PRIOR APPLICATION NUMBER: US60/322,781
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/318,184
; PRIOR FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: US60/361,663
; PRIOR FILING DATE: 2002-03-05
; PRIOR APPLICATION NUMBER: US60/396,412
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US60/322,636

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; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,817
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,816
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/323,519
; PRIOR FILING DATE: 2001-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 341
; SOFTWARE: Custom
; SEQ ID NO 152
; LENGTH: 1013
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-236-417-152

Query Match      32.3%; Score 1745; DB 12; Length 1013;
Best Local Similarity 41.7%; Pred. NO. 2.6e-149;
Matches 403; Conservative 155; Mismatches 339; Indels 70; Gaps 21;

QY 28 GVQDFVLENTSEAFIENLRERENLIYTCPLVSVNYPYEDLQIYGRHMYRG 87
Db 10 GKPDVLIDQVTIME--DFNNQLRPEKGRIRIYTYIGEVLSVNPYOELPLYGPEAIRYQG 68
QY 88 VSFYEVPPLHFAVADTVYRALRTERDQAVMISGESGAGKTEATKRLIQFYAETCPAPER 147
Db 69 RELYERPHLYAVANAAKYAMKHSRSDTCIVISGESGAGKTEASKHIMQYIAAVTNPSQR 128
QY 148 GGA--VRORLLQSNPVLBAFNAKTLRNDNSRFGKYMVDQDFKAPVGGHILSYLLEK 205
Db 129 AEVERVKDVLKSTCVLEAFAGNARTNRNHSRFGKYMVDQDFKAPVGGHILSYLLEK 188
QY 206 SRVHONHGERNFHFVYOLLEGEBETLRLGLERNPOSYLYLVKGQCAKVSINSKSDW 265
Db 189 SRVLKQHVGERNFHFVYOLLEGEBETLRLGLERNPOSYLYLVKGQCAKVSINSKSDW 247
QY 266 KMRKALSVIDFTEDEVEDLISIVASVLHGNHFADEDSNAQ-----VTENQLKYLIT 320
Db 248 QAVTEAMRVIGFSPEVESVHRILAAIHLGNIEFVETEGGLQKEGLAEEALVDHVA 307
QY 321 RLLGVEGTILREALTHRKIAKGBELLSP-INLQAAVARDALAKAVYSRTFTWLVRKIN 379
Db 308 ELTATPRDLVRLSLARTVASGGRELIEKGHTAAEASVARDACAKAVYQRLFEWVWNRIN 367
QY 380 RSLASKDAEPPSWRSTTVLGLDLYGFEVFOHNSPEQFCINVCNEKLOQLFIETLSEQ 439
Db 368 SVMPERGDRPDGDKTIVGLDLYGFEVFOHNSPEQFCINVCNEKLOQLFIETLSEQ 427
QY 440 BEYRAGTAMBPVQVFNKIIICDLVEEKFGHIIISILDECLRPGEATDLTLEKLEDTVK 499
Db 428 BEYEREGITQSVYEFNNAITVDLVERPHRGILAVLDEACSSAGTITTDRIFLQTLDMHR 487
QY 500 PHPELTHKADQKTRKSLDRG-EPRLHYAGETVSYVTGGLKNDLFPNLKXETWCSS 558
Db 488 HHLHTYSQLC--PTDKTMEGRDPRIRHYAGDVTSYVEGIDKRNDFQDFQKRLLYNS 545
QY 559 NNPIMACQF-----DKSELSPKRPETVATQFMSLLQLVILRSKEPAYIRCIKPNDAK 613
Db 546 TDPTLRAMPDQGDQDITEVT--KRPLTAGTLFKMSVALVENLASKEFFYVRCIKPNEDK 603
QY 614 QGFRPDEVLIHQVKYLGIMENLRVRAGFAYRRKYEAFLQRYKSLCPETWP--MWAGRPQ 672
Db 604 VAGKLDENHCRHQVAYLGLLENVRVRAGFASRQFYSRFLRYKMTCEYTWPNHLLGSDK 663
QY 673 DGAVLVRLHGLVKKPEYKMGRTKIFRFPKTLFATEDSLEVRQSLATKIQAAWNGFHR 732
Db 664 AAVSALLEQHGIQ--GDVAFGSKLFRSPRTLVLTQS-----R 701
QY 733 QKFLVRKRSATCQSWRGTLGRKRAAKRWAAQTIIRLIRGFIILRHSRCPENAFPLDH 792
Db 702 ARLIFI--IVULLQKAWRGTLARWF--CRLRLAIYIMWFR-----RHKVR-----HLAE 749
QY 793 VRASFLNLRLQRLPRNVLDTSWPTPPPALREASELLRELCKMKNVWVKYCRSISPEWKQOL 852
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Db 750 LQRRF--QAARQPLLYGRDLVWPLPPAVLOFPQDTCALFCRWARQVLVKNIPSDMPQI 807
QY 853 QOKAVASEIFKGGKDNY--PQSVRLFISTRLGTETISPRVLQSL-----GSEPIQYA 903
Db 808 KAKVAMGALQGLRQDWCRCRRARDYLSATDNPTASSLFAQRLKTLQDKDGFQAVLFS 867
QY 904 VPVVYDKYKGRPRROLLITPSAVIVE--DAKVQRIDYANLTGISVSSLSDSFLVL 960
Db 868 SHVRKYNR-FHKIRNRALLTDDQLHYKLDPRQYRVNRAVPELEAVTGLSVTGGDQLVVL 926
QY 961 HVQREDN 967
Db 927 HARGQDD 933

RESULT 7
US-10-080-334-164
; Sequence 164, Application US/10080334
; Publication No. US20040002584A1
; GENERAL INFORMATION:
; APPLICANT: Pena, Carol E. A.
; APPLICANT: Shimkets, Richard A
; APPLICANT: Li, Li
; APPLICANT: Shenoy, Suresh G
; APPLICANT: Kekuda, Ramesh
; APPLICANT: Spytek, Kimberly A
; APPLICANT: Vernet, Corine A. M.
; APPLICANT: Malyankar, Uriel M
; APPLICANT: Guo, Xiaojia
; APPLICANT: Gusev, Vladimir Y
; APPLICANT: Caeman, Stacie J
; APPLICANT: Boldog, Ferenc L
; APPLICANT: Furtak, Katarzyna
; APPLICANT: Tchernev, Velizar T
; APPLICANT: Patturajan, Meera
; APPLICANT: Gangolli, Esha A
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Liu, Xiaohong
; APPLICANT: Baumgartner, Jason C.
; APPLICANT: Gerlach, Valerie
; APPLICANT: Spaderna, Steven K
; APPLICANT: Zerhusen, Bryan D
; TITLE OF INVENTION: Proteins, Polynucleotides Encoding Them and Methods of
; FILE OF INVENTION: Using the Same
; FILE REFERENCE: 21402-275
; CURRENT APPLICATION NUMBER: US/10/080,334
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/270,523
; PRIOR FILING DATE: 2001-02-21
; PRIOR APPLICATION NUMBER: 60/322,712
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: 60/311,980
; PRIOR FILING DATE: 2001-08-13
; PRIOR APPLICATION NUMBER: 60/330,307
; PRIOR FILING DATE: 2001-10-18
; PRIOR APPLICATION NUMBER: 60/278,796
; PRIOR FILING DATE: 2001-03-26
; PRIOR APPLICATION NUMBER: 60/281,521
; PRIOR FILING DATE: 2001-04-04
; PRIOR APPLICATION NUMBER: 60/276,677
; PRIOR FILING DATE: 2001-03-16
; PRIOR APPLICATION NUMBER: 60/311,595
; PRIOR FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: 60/270,220
; PRIOR FILING DATE: 2001-02-21
; PRIOR APPLICATION NUMBER: 60/274,295
; PRIOR FILING DATE: 2001-03-08
; PRIOR APPLICATION NUMBER: 60/318,526
; PRIOR FILING DATE: 2001-09-10
; PRIOR APPLICATION NUMBER: 60/286,548
; PRIOR FILING DATE: 2001-04-25
; PRIOR APPLICATION NUMBER: 60/291,765
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; PRIOR FILING DATE: 2001-05-17
; PRIOR APPLICATION NUMBER: 60/270,797
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/276,400
; PRIOR FILING DATE: 2001-03-16
; PRIOR APPLICATION NUMBER: 60/270,810
; PRIOR FILING DATE: 2001-02-23
; NUMBER OF SEQ ID NOS: 388
; SOFTWARE: Patent In Ver. 2.1
; SEQ ID NO 164
; LENGTH: 688
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-080-334-164

Query Match      32.1%; Score 1732; DB 15; Length 688;
Best Local Similarity 50.0%; Pred. No. 2e-148;
Matches 349; Conservative 132; Mismatches 187; Indels 30; Gaps 10;

Qy 28 GVQDFVILENTSAAFTENLRFRNLIYIYIGPVLSVNPYRDQIYSRQHMERYRG 87
Db 7 GVEDLVLL-TYLNPAVLHNLKRYLDLIYIYIGLVSVNPYKQLEPIYTPYVKKYRG 65
Qy 88 VSFVEVPHLPAVADTYVRLALTRERDOAVVMISGESGAGKTEATKRLQLQFYAETCPAPER 147
Db 66 KSGELPPEVFAIADNAYRNMLNDKQNSIIISGESGAGKTEATKTMQVLAASVSGSGS 125
Qy 148 GGAVERDLLOSNPVLEAFNAKTLNRDNRFRGMYDMVQDFKCAPYGGHLSVLLKSR 207
Db 126 VGSVEDQILSNPFLFAFGNAKTLNRNNSRFGFIEIHFDAKGIYGAKIETVLLKSR 185
Qy 208 VHQNHGERNPHVYQLLEGSEETLRLGLERNPQSYLYLVKQCAKVSINDKSDWKV 267
Db 186 VWSQAGERNYHIFYQLLAGASELKKGLKSPEDYRVLNQGGCLTVGIDDAEFKE 245
Qy 268 MRKALSVIDTEDEVEDLLSVASVLHGNHFAEDSDNAQVTEN--QKYLTRLGV 325
Db 246 TLNAMRVIGFSEBEESESIFKILAAIHLGNIEPEEGENDNAAETVKDKEELDNAELLGV 305
Qy 326 EGTTLREALTHRKIIAKGEBLLPLNLEQAAAYARDALAKAVYSRTFTVLVYKKNRSIASK 385
Db 306 DPPELEKALTAKTKTGGEVITVPLTVEQALDARDALAKAIYSLFDFWLVRINQSLSFK 365
Qy 386 -DAESPSWRSTTVGLLDIYGVFEVQHNSEQFCINCYNEKLOQLFIELTLKSEQBEYEA 444
Db 366 RDG-----STNFTGLVDIYGFIFEFKNSPEQLCINYANEKLAQFFNQHVFKLEQEYER 419
Qy 445 EGIAWEPVQVFNKKIICDLVEEKFGIISILDECLRPGEATDILTEKL-EDTVKPHPH 503
Db 420 EGIWTFIDFDQDCIDILIEKKPLGLSLDLDECRFP--KGTDTQTEKLNKQHLKHPH 478
Qy 504 FLTHKLADQKTRKSLDRGEFRLHYAGEVTVSYTGFLDKNDLILFNLKETMCSMNPI 563
Db 479 F-----SKPKN-GRTEPIIKHYAGDVTVYTGFLKKNKOTLSDDLIELLQSSKNPLI 530
Qy 564 AQCFDK-----SELSDKKPEVATQVQKMSLLQLVELRSKEPAYIRCIPKNDKQ 614
Db 531 ALLFPPEAGQTSSAPKESKAKKFFQTVGSQFESLNEMLDTLNTNPHFIRCIPKNEEK 590
Qy 615 PGRFDEVILRHQVKYLGMLNLRVRAFGAYRKYEAFLQRYKSLCPETWPMWAGRPQG 674
Db 591 PGDFDSSLVLHQYUIGVLETHIRAGFPYRPLPFDFELQRYRVLLPDTWPMWGGDAKEA 650
Qy 675 VAVLRHLGYKPEYKMGRTKIFIRFPKTLFATEDSL 712
Db 651 CELLSQGLGDEDEYQIGTKTVFLR-PGQLAELEELRE 687

RESULT 8
US-10-336-472-126
; Sequence 126, Application US/10336472
; Publication No. US20040043929A1
; GENERAL INFORMATION:
```


Db 429 KQOEYEREGITWQSVYFNNAITVDLVERPHRGILAVLDEACSSAGTITDRIFLQTL 488
 Qy 496 DTVPHPHFLHKLADQKTRSLDRG--EPRLLHAGVYTVYVGTGFLDKNDLLIFRNLIKET 554
 Db 489 THRHHLHYTSRQLC--PTDIMEFGRDFRKHVAGDVTVSVGEFIDKNDRLDFQDFKRL 546
 Qy 555 MCSNNPIMACQF-----DKSELSDDKKRPETVATQFKMSLLQLVEILRSKEPAYIRCIKP 609
 Db 547 LYNSTDPTRLRAMPDGOQDITEVT--KEPLTAGTLFKNSMVALVENLASKEFFVYRICKP 604
 Qy 610 NDAKOPGRFDEVILTRHQVKYLGILMENLVRVRAGFAYRRKYEAFLORYKSLCPETWP--MWA 668
 Db 605 NEDKAVAGKLDENHCHQVAYILGLENVRVRAGFASRQPYSRFLRYKMTCEYTPNHL 664
 Qy 669 GRPDQGVAVLVRHLGYKPEEYKMGRTKIFIRFPKTLFATEDSLEVRQSLATKIQAAWRG 728
 Db 665 GSDKAASALLEQHLQ--GDVAFGHSKLFIRSPRLVTLFOS----- 705
 Qy 729 FHMROKFLVRKSAICIQSWWRGTLGRKKAARKWAAQTIRRLIRGFILRHSRCPENAF 788
 Db 706 ---EARLIP--IVLLQKAWRGTLARWR--CRLRLAITYTIMRWR-----RHKYRA----- 750
 Qy 789 FLDHVRASEFLNLRLRQLPVNLDSWTPPPALREASELLRELCKMKNWVKYCRSISPEW 848
 Db 751 HLAELQRRF--QAARQPLYGRDLVPLPAVLQFPQDTCALFCRWRARQLVKNIPPSD 808
 Qy 849 KQQLQOKKAVASEIPKGGKONY--PQSVRLFISTRLCTEISPRVLOS-----GSEP 899
 Db 809 MPQIKAKVAANGALQGLRQDWGCRARAWDYLSATNPASSLFAORLTKLKDQGFGA 868
 Qy 900 IQYAVPVVYKRGYKPRQLLTPSAVIVE---DAKVORIDYANLTAIGSVSSLSDS 956
 Db 869 VLFSSHVYKVNRR--PHKIRNALLITDQHLKLDPRQYVRMRAVPLEAVTGLSVTSGDQ 927
 Qy 957 LFLVHLVQREDN 967
 Db 928 LVVLHARGQDD 938

RESULT 10
 US-10-104-047-3280
 ; Sequence 3280, Application US/10104047
 ; Publication No. US20030236392A1
 ; GENERAL INFORMATION:
 ; APPLICANT: HELIX RESEARCH INSTITUTE
 ; TITLE OF INVENTION: No. US20030236392A1e1 full length cDNA
 ; FILE REFERENCE: H1-A0105
 ; CURRENT APPLICATION NUMBER: US/10/104,047
 ; CURRENT FILING DATE: 2002-03-25
 ; PRIOR APPLICATION NUMBER:
 ; PRIOR FILING DATE:
 ; NUMBER OF SEQ ID NOS: 4096
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO 3280
 ; LENGTH: 1098
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-104-047-3280

Query Match 31.9%; Score 1722.5; DB 15; Length 1098;
 Best Local Similarity 38.2%; Pred. No. 3.3e-147;
 Matches 401; Conservative 152; Mismatches 303; Indels 193; Gaps 23;

Qy 28 GVQDFVLEENFTSEAFIENLRFRFENLTYTIGVPLVSVNPNYRDLQIYRQHMERYRG 87
 Db 18 GVDDWVLLPOIT--EDAIAENLRFRFMDYITFTIGSVLISVNPFKQMPYFTDREIDLYQG 76
 Qy 88 VSYFVPPHFLFAVDATYRALRTERDDQAYMISGESGAGKTEATKRLQFYAETCPAPER 147
 Db 77 AAQYENPPHYALTNDMYRNWMLDNCQCVIIISGESGAGTKVAKYINGVISKVSGGK 136
 Qy 148 GGAVRRLQSNPVLFAFGNAKTLRNDNSRFRQKYMVDQFDFKGAIPVGGHILSYLLEKSR 207

Db 137 VQHVKDIILOSPLLEAFNAKIVRNNSRFRGKYFEIQSRGGEPPGGKISNELLBKSR 196
 Qy 208 VVQNHGERNPHFYQYLLGEBEETLRLGLERNPQSYLYLVKQCCAKVSINDSKDWKV 267
 Db 197 VVQNEENRNFYIYQLLEGASQEQRLNGL--WTFDYVYVYNQSDTVQVQDTRDSRGE 255
 Qy 268 MRKALSVIDTDEVEDLLSIVASVLHNLHINHADEDSN--AQVTENQLKYLTRLIGVE 326
 Db 256 TLSVMQVIGIPPSIQQLVLQVAGILHGNISFC--EDGNVYARVESVLLAFAYLLGID 313
 Qy 327 GTTLREALTHRKIIA-----KGEELLSPNLNQAAAYARDALAKAVYSRTETWLVRKINRSL 382
 Db 314 SGRLOEKLTRKMSRWGRGSEINVTLVNVEQAAVTRDALAKGLYALFLDFLVRAINRAM 373
 Qy 383 ASKDAASPSWRSTVTLGLLDIYGFVFOHNSFFQFCINYCNEKILQQLFIBLTAKSEOEY 442
 Db 374 -QKPQBEYS-----IGVLDIYGFIFQKNGPFOFCINFVNEKILQQLFIBLTAKSEOEY 426
 Qy 443 EASGLAWEPOVYFNKKIICDLVEEKK--GIISILDEECI--RPGEATDLTFLEKLEDT 497
 Db 427 VQGIIRWTPIQYFNKVVCDLIENKLSPPGIMSVLDDVCAVTHATGGADOTLLKQLQAA 486
 Qy 498 VKPHPHFLTHKLADQKTRKSLDRGEFRLHYAGEVTVSVTGFLLDKNDLLFRNLKETMCS 557
 Db 487 VGTHEHF-----NSWSAG--FVHHYACKVSYDVSGFCERRDVLFSLLIELMQT 534
 Qy 558 SNNPIMACQFCKSELSDDK--RPTVATQFMSLLQLVEILRSKEPAYIRCIKPNDAKQPG 616
 Db 535 SEQAFRLMFLPEKLDGDKGRPSTAGSKIKQANDLVATLMRCTPHYIRCIKNETKPR 594
 Qy 617 RPEVLRIRHQVKYLGILMENLVRVRAGFAYRRKYEAFLORYKSLCPETWPNWAGPDQVA 676
 Db 595 DWENRKKQVEYLGUKENIRVRAGFAYRRKYEAFLORYKSLCPETWPNWAGPDQVA 654
 Qy 677 VLVRHLGYKPEEYKMGRTKIFIRFPKTLFATEDSLEVRQSLATKIQAAWRGPHWRQKPL 736
 Db 655 HLLRAVNMPEPQYQMGSTKVFKNPESLFLLEBVRERKEDGFPARTIQKAW----- 704
 Qy 737 RVKRSALCISQSWWRGTLGRKKAARKWAAQTIRRLIRGFILRHSRCPENAFFLDHVRAS 796
 Db 705 -----RRHVAVRKY----- 713
 Qy 797 FLNLRLRQLPRNVLDTSWTPPPALREASELLRELCKMKNWVKYCRSISPEWQQLQOKA 856
 Db 714 -----EEMRE----- 718
 Qy 857 VASEIFKGGKDNYPQSVPRFLFISTRIGTEISPRVLQSLGS--EPIQYAVPVVYKRGYK 915
 Db 719 EASNILLNKKERRRNSINRNFVGDYLGEE--RPELRQFLGKREVDFAVSVKYDRR--FK 776
 Qy 916 PRPRQLLTPSAVIVVEDAKV-----KORIDYANITGISVSLSDSLFVLHV 962
 Db 777 PIKRDILTPKCVYVIGREKVKKPGQGVCEVKKKVDIQALRGVSLSTRQDDFFIL-- 834
 Qy 963 QREDNKKQGDVVVLQSDHVTETLTKT---ALSADRVNNININQGSITFA----- 1007
 Db 835 -QED-----AADSFLSVFKTFEVSLLCKRFEATRRLPLFTFSDTLQFRVKKEGW 884
 Qy 1008 GQPGRDIIDFTSGSELLITKAKNGHAY 1036
 Db 885 GGGGTRSVTSRFGPGLAVLVKVGGRITLV 913

RESULT 11

US-10-202-481-2

; Sequence 2, Application US/10202481

; Publication No. US20040018567A1

; GENERAL INFORMATION:

; APPLICANT: Vallone, Marcy K

; APPLICANT: Wong, Brian R

; APPLICANT: Masuda, Estaban

; APPLICANT: Powell, Mark

; TITLE OF INVENTION: Modulators of B-lymphocyte Activation, Myosin-1F Compositions ar

```

; TITLE OF INVENTION: cf Use
; FILE REFERENCE: A-71312/RMS/TAL/DHR
; CURRENT APPLICATION NUMBER: US/10/202,481
; CURRENT FILING DATE: 2002-07-23
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 2
; LENGTH: 1098
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-10-202-481-2

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Query Match 31.8%; Score 1714.5; DB 15; Length 1098;
Best Local Similarity 38.1%; Pred. No. 1.8e-146;
Matches 400; Conservative 152; Mismatches 304; Indels 193; Gaps 23;

QY 28 GVQDFVLLNFSTSEAAFIENRRRENLIYTGIVLVSVNYPYEDLIQYSRQHMERYRG 87
DB 18 GVUDDVLLPQIT-EDAIANLRKRFDYFIYIGSVLISVNPFPQMPYFFTDREIDYQG 76
QY 88 VSFYEPHPFLAVADVTVYRALTRERDQAVMLISGSGAGKTEATKRLIQFYAETCPAPER 147
DB 77 AAQYENPPHYALTNDWYERNMLDCENQCVIISGSGAGKTVAAKYINGYISKVSGGGEK 136
QY 148 GGAVRDLQSNPVLFAFNATKLRNDSRRGKYMVDQFQKAPVGGHLSYLLLEKSR 207
DB 137 VQHVKDIIILQSNPLLEAFNAKTVNNNSRRGKYFEIQFSGGEPDGGKISLLEKSR 196
QY 208 VVHNGEGERNEHVYQLLEGGEEFTLRGLERNQSYLYLVKGQCAKVSINDKSWKV 267
DB 197 VVQCNERNRHHYQLLEGAGEQORNLGL-MTPDYIYINQSDTVQVDDTDDRSDGE 255
QY 268 MRKALSVIDFTEDEVEDLLISIVASVHLHGNHFAADEDSN-AQVTENOLKYLTRLGLVE 326
DB 256 TLSAMQVIGPISIQQLVQLVAGILHLGNTSFC--EDGNVAVRESVDLLAFAYLLGID 313
QY 327 GTTLREALTHRKIIA-----KGELISPLNLEQAAVARDALAKAVYSRTFTWLVRKINRSL 362
DB 314 SGRLOEKLSRRKMSRWGGRSSEINVLNVEQAAVTRDALAKGLYARLFDPLVEAINRAM 373
QY 383 ASKDAESPMSRTTVLGLDIYCFEVQHNSEFQRCINYEKLOQLFIETLKSEOEY 442
DB 374 -QKQOEYS-----IGVLDIYGFIFQNGEQQFCINFEKLOQFIETLKAEOEY 426
QY 443 EAGTAWEPVQVFNKKIICDLVEERFK--GIISILDECL--RPGEATDLTFLEKLEDT 497
DB 427 VQEGIRWTIYQFNKKVCDLLENKLSPPGIMSVLDDVCATMHTATGGADOTLLQKQAA 486
QY 498 VKPHPHLTHKLADQKTKSLDRGFRLLHYAGEVTVSVTGFLDKNNDLFLRNLKETMCS 557
DB 487 VGTHEHF-----NSWSAG-FVTHHYAGKVSVDVSGFCERNRDLVFSDLIELMQT 534
QY 558 SNNPIMAQCFDKSELSDKK-RPETVATQFKMSLLQLVEILRSKEPAYIRCIKNDKQPG 616
DB 535 SEQAFURMLFPBKLDGDKRGRESTAGSKIKQANDVATLMRCPTPHYIRCIKNETKHAR 594
QY 617 RFDEVLIRHQVKYGLIMENLRVRRAGFAYRRKYEAFLQRYKSLCPETWPMWAGRPDQGA 676
DB 595 DWEENRVKHQVYGLKENIRVRRAGFAYRRQYQPAFLQRYAILTPETWPRWGRDQGVQ 654
QY 677 VLVRHLYGKPEYKMGRTKIFIRPKTLFATEDSLSEVRQSLATKIQAAMRGFWHRQKFL 736
DB 655 HLLRAVNMEPDQYQMGSTKVFVKNPESLFLLEVEERKFDGFARTIQAM----- 704
QY 737 RVKRSALICQSWRGTLGRKAAKPKWAAQTIRRLIRGFLRHSRCPENAFLLDHVRAS 796
DB 705 -----RRHVAVRKY----- 713
QY 797 FLINLRQLPRNVLDTSWTPPPALREASELLRELCKMNMVWKYCRSISPEWKOQLQKA 856
DB 714 -----EMRE----- 718
QY 857 VASEIFGKKDNPQSVPRFLFSTRGLTEISPRVLQSLG-SEPIQYAVPVVYDKRGYK 915

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; PRIOR APPLICATION NUMBER: 10/055,569
; PRIOR FILING DATE: 2001-10-26
; PRIOR APPLICATION NUMBER: 10/080,334
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 10/092,900
; PRIOR FILING DATE: 2002-03-07
; PRIOR APPLICATION NUMBER: 10/136,826
; PRIOR FILING DATE: 2002-05-01
; PRIOR APPLICATION NUMBER: 10/236,417
; PRIOR FILING DATE: 2002-09-06
; PRIOR APPLICATION NUMBER: 60/345,092
; PRIOR FILING DATE: 2002-01-04
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 230
; SOFTWARE: CuraseqList version 0.1
; SEQ ID NO 26
; LENGTH: 1096
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-336-472-26

Query Match      31.7%; Score 1709.5; DB 12; Length 1096;
Best Local Similarity 38.2%; Pred. No. 5.1e-146;
Matches 400; Conservative 148; Mismatches 307; Indels 193; Gaps 23;

QY      28  GVQDFVILENTSEAAFTENLRPRFRENLIITYIGVLVSVNPRVDLQIYGRQMERYG 87
Db      18  GVDWVLFPQT-EDAIANLRKPRMDDYIFTYIGSVLISVNPFKQMPYFTDREIDL YQG 76

QY      98  VSFYEVPHLFAVADTVYRALTRBRRDQAVMISGESGAGKTEATKRLQFYAETCPAPER 147
Db      77  AVQYENPHIYALTDMYRNMLDCENQCVLISESGAGKTVAAKIMGYISKVSGGGEK 136

QY      148  GGAVDRLLQSNPVLFAFGNAKTLRNDNNSRFGKMYDQDFKGAPVGGHTLSVILEKSR 207
Db      137  VQHVKDITLQSNPLLEAFGNAKTVRNNSSRFGKYFEIQFSRGGEPEGDKISINFLLEKSR 196

QY      208  VVHQNHGRNPHVYVLLEGEEETLRLGHERNPQSVLYLVKQCAKVSINDKSWKV 267
Db      197  VVMQNEHNRPHIYYQLLEGASQQRQNLGL-MTFDYIYYLNQSDTYQVGDGTDSDSDFGE 255

QY      268  MRKALSVIDFTEDEVEDLLSIVASVLHGNTHFAADESNAAQVTTENQLKYLTELLGVEG 327
Db      256  TLSAMQVIGIPPSIQQLVLQVLVAGILHGNISFC--EDGNVARVESVDLAPFVLLGIDS 313

QY      328  TTIREALTHKRIIA----KGELLSPINLEQAAVARDALAKAVYSRTFTWLVRKINRSLA 383
Db      314  GRQEKLTSRKMDSRWGRSGRSESINVTLNVEQAAVTRDALAKGLYARLDFDLVEAINRAM- 372

QY      384  SKDAEPPSWRSTTVLGLDLYIGREVFQHNSEQFCINCYNEKLOOLFIETLTKASEQEYE 443
Db      373  QKQEEYS-----IGVDLYIGFEIFQNGFEQFCINPVNEKUQOIFIELTKRAEQEYV 426

QY      444  AEGIANPQVYFNKKIICDIVEEKFK--GIISILDERCL--RPGEATDITLFEKLEDTV 498
Db      427  QEGIRWTPIQVFNKKVVCDLIENKLSPPGIMSVLDDVCATMHAATGGADQTLQKLAQAV 486

QY      499  KPHEPFLTHKLADQKTRKSLDRGEFRLLHYAGEVTVYVTFGLKNNDLLFRNLKBTWCSS 558
Db      487  GTHEHF-----NSWSAG-FVTHYVAGKSYDVSGFCERNRDLVFSDLIELMQTS 534

QY      559  MNPTMAQCDFKSELSDKK-RPETVATQFKMSLLQLVEILRSKBPAYTRCIPKNDAKOPGR 617
Db      535  -EQFLRMLFPEKLDGDKGRPSTAGSKIKQANDLVATLMRCTPHYIRLCIKPNETKRPRD 593

QY      618  FDEVLRHVKVYIGLMENLRVRAGFAYRRKRYEAFLQRYKSLCPETWPMWAGRPQGVAV 677
Db      594  WEENRVKHQVEYLGLENIRVRAGFAYRRQFAKFLQRYAILPETWPRWRGDERQGVQH 653

QY      678  LVRLHGLKPBEEYKMGRTKIFIRPKTLFATEDLSLEVRQSLATKIQAWRGFWHQKFLR 737
Db      654  LLRAVNMEPOYQMGSTKVPKPSLLEEYRERKFRDGFARTIQAW----- 702

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QY 738 VRSACIQTQSWRGTLGRKKAAGKAAQATIRRLIRGFIILHSPRCPENAFFLDHVRASF 797
Db 703 -----RRHVAVRKY-----711
QY 798 LLNLRLRQLPRNVLDTSWPTPPALREASELLRELCKMVMWKYCRSISPEWKQOLOQKAV 857
Db 712 -----EMRE-----E 717
QY 858 ASEIFKGKKNYPQSPRLFTSLRGTDEEISPRVLQSLGS-EPIQVAVPVVKYDRKGYP 916
Db 718 ASNILLNKERRRNSINRNFVGYLGLLE-RPELRQLGRVRVDFADSVTKYDRR-FKP 775
QY 917 RPROLLITPSXAVIVEDAKV-----KORIDYANLTGIVSSLSLSDSLFVILHVQ 963
Db 776 IKROLILLTPKCVYVIGREKVKKGPEKQGVCEVLKKKVDIQALRGVSLSTEQDDFFIL--- 832
QY 964 REDNKQKGDVVQLQSDHVIETLTKT--ALSADRVNNININQSGITFA-----G 1008
Db 833 QED-----AADSPLEGSVFKEFVSLCKFEATERRPLFTSDRLQFRVKKEGWG 883
QY 1009 GPGRGDIIDFTSGSELLITKAKNGHLAV 1036
Db 884 GGGRTSVTFSGFGCDLAVLKVGGRTLV 911

RESULT 13
US-10-236-417-66
; Sequence 66, Application US10236417
; Publication No. US20040048256A1
; GENERAL INFORMATION:
; APPLICANT: Agee et al.
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-442C
; CURRENT APPLICATION NUMBER: US/10/236,417
; CURRENT FILING DATE: 2003-01-06
; PRIOR APPLICATION NUMBER: US60/318,120
; PRIOR FILING DATE: 2001-09-01
; PRIOR APPLICATION NUMBER: US60/318,430
; PRIOR FILING DATE: 2001-09-10
; PRIOR APPLICATION NUMBER: US60/322,781
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/318,184
; PRIOR FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: US60/361,663
; PRIOR FILING DATE: 2002-03-05
; PRIOR APPLICATION NUMBER: US60/396,412
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US60/322,636
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,817
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/322,816
; PRIOR FILING DATE: 2001-09-17
; PRIOR APPLICATION NUMBER: US60/323,519
; PRIOR FILING DATE: 2001-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 341
; SOFTWARE: Custom
; SEQ ID NO 66
; LENGTH: 1096
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-236-417-66

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Query Match      31.7%; Score 1709.5; DB 12; Length 1096;
Best Local Similarity 38.2%; Pred. No. 5,le-146;
Matches 400; Conservative 148; Mismatches 307; Indels 193; Gaps 23;
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Qy	28	GVOFVLENTSEAFIENLRFRFNLTYYIGVLVSVPYRDIQIYSROHMEYRG	87
		: : : :	
Dd	18	GVDDMWLLPQIT-EDAIANLRKRFWDITYFTIGSVLISVNFPKMPYFTDREIDLYQG	76
		: : : :	
Qy	88	VSYEYVPPHLFAVDVTYRALFTRERDQAVMISGESGAGKTETAKELLQFYAETCPAPER	147

Db 1 VEDMAEL-TYLNPSVLHNLKRYKADLIYTYSGVLVSVNPNYKRLPIYTBEEFIKVRGK 59
 Qy 89 SFYVPPHFAVATVYRALKTERDOAVMISGSGAGKTEATKRLQFVAETCPAPERG 148
 Db 60 RRYELPHIFAIADEAYESMLSDKENSILISGSGAGKTEATKRVMOYLA----AVSGG 115
 Qy 149 -----GAVDRLLQSNPVLEAFNGNAKTLRDNNSRFGKYMVDQDFKGAAPVGGHLSYLL 203
 Db 116 NGKVGVRVEDQILQSNPILFAFGNAKTRNNNSRFGKYIEIQDKTKIVAGAKIENYLL 175
 Qy 204 EKSRRVHQNCHERNFHFVYQLLEGSEBETLRLGLERNPQSYLYLVKQCAKVSINDKS 263
 Db 176 EKSRRVYQTPGERNFHIFYQLLAGASQQLKELNL-TDPDDYHVLNQGCGYTVDDGIDSE 234
 Qy 264 DWKMRKALSVIDFTEDEVEDLLSIVASVLHNLGNIHF-----AADSDSNAQVTTENQL 316
 Db 235 EFETDKAMDLTGFSDDEEQLSIFRVAAILHLGNIKFKYQRKKEAABDDT-----KAL 288
 Qy 317 KYLTRLLGVEGTTUREALTHRKIIITAKGEELSPNLQAAAYARDALAKAVYSRTFTWLVR 376
 Db 289 QIAAELLGVDKALEKALLSRRIKGTGEGVTPQNVQANAYARDALAKALYSRLFDWIVN 348
 Qy 377 KINRSKASDAESPSWSTTVLGLDIYGEVFGHNSFQFCINYNCKEQLQOLFIELTLK 436
 Db 349 RINKSLDFKAKEGANF-----IGVLDIYGEIFEKNSFEQFCINYNCKEQLQOLFENHMFK 403
 Qy 437 SEQEYEAEGIAEPVQYFNKKIICDLVEBKFGIISILDEECILRPOEATDLTFLKLED 496
 Db 404 LEQEYKREGIEWTFIDFGNQPCIDLIIEKPPGILSLDBECRFP-KATDQFELDLYS 462
 Qy 497 TVKXPHFLTHKLADQKTRSLDRGEPRLHYAGEVTVYTGFLDKNDLLFRNLKETMC 556
 Db 463 EFSNHPHFKPRQKKS-----FIKHVAGDVEYNVEGELEKKNKDLFDLIELLK 514
 Qy 557 SSNNPMAOCF-----DKSELSKKRPE-----TVATOPKMSLLQLVLILSKEPAYI 604
 Db 515 SSNNPILAEFPDYEEADPSLSKKRKITTKSNFIVTGAQFKESLNTLMTLSSTNPHFV 574
 Qy 605 RCIKPNDKAPGRFDEVLIRHVKYLGMLMENLRVRAGFAIRRYKRYEAFIORYKSLCPETW 664
 Db 575 RCIKPNEKKPGVFDASVLHQLRCLGVLSGIRIRAGFSPRITPFEFLQRYILAPKTW 634
 Qy 665 PMWAGRPQDGA-VLVHRLGYKPEEYKMGRTKIFIR 699
 Db 635 PKWSGAKKAGACELLQALNLDKEEYQFGTKIFFR 670

RESULT 15
 US-10-369-493-5164
 ; Sequence 5164, Application US/10369493
 ; Publication No. US20030233675A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Cao, Yongwei
 ; APPLICANT: Hinkle, Gregory J.
 ; APPLICANT: Slater, Steven C.
 ; APPLICANT: Goldman, Barry S.
 ; APPLICANT: Chen, Xianfeng
 ; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
 ; FILE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
 ; FILE REFERENCE: 38-10(52052)B
 ; CURRENT APPLICATION NUMBER: US/10/369,493
 ; CURRENT FILING DATE: 2003-02-28
 ; PRIOR APPLICATION NUMBER: US 60/360,039
 ; PRIOR FILING DATE: 2002-02-21
 ; NUMBER OF SEQ ID NOS: 47374
 ; SEQ ID NO 5164
 ; LENGTH: 1100
 ; TYPE: PRT
 ; ORGANISM: Caenorhabditis elegans
 US-10-369-493-5164

Query Match 29.0%; Score 1564; DB 15; Length 1100;
 Best Local Similarity 34.7%; Pred. No. 9.7e-133;

Matches 367; Conservative 172; Mismatches 320; Indels 200; Gaps 21;
 Qy 27 VGVODFVLLNFTSEAAFIENLRFRPRENLIYTYGFLVSVNPNRYDLQIYSRQHMERYR 86
 Db 14 VGDDMDVLLPKLT-EQSIENLKKRLQANSIFTYIGFVLVSVNPFKQMPYFTEKEMLLYQ 72
 Qy 87 GVSFEYEPHLEAVADTVYRALKTERDOAVMISGSGAGKTEATKRLQFVAETCPAPE 146
 Db 73 GAQYENAPHIYALADNMVYENMLINDNESQCVILISGSGAGKTVNAKIFIMYISISGGQ 132
 Qy 147 RGGAVDRLLQSNPVLEAFNGNAKTLRDNNSRFGKYMVDQDFKGAAPVGGHLSYLLKS 206
 Db 133 KVQHILQSNPILFAFGNAKTRNNNSRFGKYIEIVFSRGGEPIGGKLSNFLLEKS 192
 Qy 207 RVYHQNCHERNFHFVYQLLEGSEBETLRLGLERNPQSYLYLVKQCAKVSINDKS 266
 Db 193 RVYHQNCHERNFHFVYQLLEGSEBETLRLGLERNPQSYLYLVKQCAKVSINDKS 251
 Qy 267 VMKALSVIDFTEDEVEDLLSIVASVLHNLGNIHF-----AADSDSNAQVTTENQL 326
 Db 252 STLHMKVGVNDQDQLEVLIRIVATVILHIGNITF-TEENNFAAVSGKYLEYPAFLGLT 310
 Qy 327 GTTLREALTHRKIIIAK-----GEELSLPLNLEQAAAYARDALAKAVYSRTFTWLVR 382
 Db 311 SADIEAKLTGRKMESKMGTKQKEIDMKLVNQASYYTDAWVKAIYARLFDYLVKYNDA 370
 Qy 383 ASKDAESPWSRSTTVLGLDIYGEVFGHNSFQFCINYNCKEQLQOLFIELTLKSQEEY 442
 Db 371 ---NITSQTSNDNFSGVILDIYGEIFEKNSFEQFCINYNCKEQLQOLFIELTLKSQEEY 427
 Qy 443 EABGIAWEFPVQYFNKKIICDLVEEKF-KGIISILDEECILR-----PGATDLTLEKLED 498
 Db 428 VREGINKWTEIDYFNKIVCDLIETKPPGIMSLDDTCAQNHGQRGVDRQLTTLSKSF 487
 Qy 499 KPHPHFLTHKLADQKTRSLDRGEPRLHYAGEVTVYTGFLDKNDLLFRNLKETMCSS 558
 Db 488 AGHPHE-----GPGSDSFVIKHVAGDVTYVDFCDNRNLDVLYPDILLMQKS 535
 Qy 559 MPIMAAQCFDKS-ELSDKXRPETVATOFKMSLLQVLVEILRSKEPAVIRCIKPNDAKOPGR 617
 Db 536 SRPFFQALFPENVAASAGKRPFTTFTKIRTOANTLVESLMKSGSPHYVRCIKNETKRPND 595
 Qy 618 FDEVLIRHVKYLGMLMENLRVRAGFAIRRYKRYEAFIORYKSLCPETWPMWAGRPQDGA 677
 Db 596 WEESRVKHQVHVLGLENIRVRAGFAIRRYKRYEAFIORYKSLCPETWPMWAGRPQDGA 655
 Qy 678 LVRHLGYKPEEYKMGRTKIFIRFPKTFEATEDSLEVRQSLATKIQAAWRFHWKQFLR 737
 Db 656 ICDSVHMEKNQYOMGKTKLIFVKNPESLFLLEETREKRPDGYARVIOKAMRQFSAR---- 710
 Qy 738 VKESAICIQSWWRGTLGRKKAARKWAQTIIRRLINGFILRHSRCPENAFDLHVRAF 797
 Db 711 ----- 710
 Qy 798 LILNRLRQLPRNVLDTSWPTPPPALREAGSELLRELQCMQNMVWKYCRSISPEWKQQLQKAV 857
 Db 711 -----KOHIKQKQEQ 719
 Qy 858 ASEIFPKGKNYQSPVRLFLSTRIGTEISPRVLQSL--GSEPIQYAVPVVYKDRKGYK 915
 Db 720 AADLMVGKERRRYSLNRFNFGVYIGLEH--HPTQLSLVGKQRVLFACTANKYDRK-PR 776
 Qy 916 PRPRQLLTTPSAVILVEDAKV-----KQRTDYANLTGISVSSLSDSFLVLHV 962
 Db 777 VTKLDDLLTVNHLTLIGKEKVNKGPEKGIKEVIRKQFQDLPOIKSIGLSPYQDDFVILYL 836
 Qy 963 QREDNKQKQGVVLQSDHVIETLTK-----TALS---ADRVNN-----ININQSGI 1004
 Db 837 GNDDYSS-----LLETFKTEFTALS KAYKERTNGTLHLDLFRSSHVVSYKKMKF 886
 Qy 1005 TFAGGP-----GRDGIIDFTSGSELLITKAKNGLHAVVA 1038
 Db 887 DFDGKRTVQFGNDG-----TSSAEK--TLKPNKGLVNS 919

Search completed: March 30, 2004, 15:21:46
Job time : 82.8075 secs

Result No.	Score	Query		DB	ID	Description
		Match	%			
1	1191	22.1	1939	4	US-09-310-187A-1	Sequence 1, Appli
2	1190	22.0	1932	4	US-08-875-435B-4	Sequence 4, Appli
3	1169.5	21.7	1972	4	US-08-875-435B-3	Sequence 3, Appli
4	1150.5	21.3	1285	4	US-09-976-594-507	Sequence 507, App
5	1146	21.2	1120	4	US-09-147-404-1	Sequence 1, Appli
6	1105	20.5	2548	4	US-09-172-432-1	Sequence 1, Appli
7	1078.5	20.0	1236	4	US-09-883-134-4	Sequence 3, Appli
8	1018	18.9	1886	4	US-08-938-105-3	Sequence 4, Appli
9	575.5	10.7	1695	4	US-09-866-108A-15753	Sequence 3, Appli
10	519.5	9.6	1581	4	US-09-866-108A-15754	Sequence 15754, A
11	497	9.2	2568	4	US-09-866-108A-3	Sequence 3, Appli
12	160	3.0	1809	3	US-09-012-515A-12	Sequence 12, Appl
13	160	3.0	1809	3	US-08-360-144A-12	Sequence 12, Appl
14	160	3.0	1809	3	US-09-012-504A-12	Sequence 12, Appl
15	160	3.0	1809	4	US-09-012-399A-12	Sequence 12, Appl
16	160	3.0	2549	4	US-08-471-112A-3	Sequence 3, Appli
17	160	3.0	2549	5	PCT-US95-06722-12	Sequence 12, Appl
18	154	2.9	2549	4	US-08-265-967C-1	Sequence 1, Appli
19	154	2.9	2549	4	US-08-305-790B-2	Sequence 2, Appli
20	152.5	2.8	103	3	US-08-305-223-395	Sequence 395, App
21	132.5	2.5	3878	4	US-09-914-259-11	Sequence 11, Appl
22	130	2.4	1529	4	US-09-134-001C-3945	Sequence 3945, Ap
23	126	2.3	567	4	US-09-134-001C-3762	Sequence 3762, Ap
24	124	2.3	1093	5	PCT-US93-03077-1	Sequence 1, Appli
25	123.5	2.3	1333	3	US-09-356-952-2	Sequence 2, Appli
26	123.5	2.3	1333	4	US-09-976-594-312	Sequence 312, App
27	122.5	2.3	1319	2	US-08-290-731C-2	Sequence 2, Appli

Db 406 KVGNEYVTKQSVQVYVYSGALAKAVYKMFNMWVTRINATLETKPQ-----YFIG 459
QY 400 LLDLYGFEVQHSFEQFCINCYCNKELQQLFIELTKSEBEBYAEAGIANEPVOYFNKI 459
Db 460 VLDIAGEFIEDFNFEQFCINFTNEKLCQFPNHHMFVLEQEBYKKEGIEWTFIDFGMDLQ 519
QY 460 IC-DLVEBEKFGIISIDEECLRGEATDITLFEKLEDVTKPHPHFTLTKLADQKTR--K 516
Db 520 ACIDLI-EKPMGINSILEECMFP-KATDMTFKALYD-----NHLGKSNQKPNV 571
QY 517 SLDRGFRLLHYAGEVYTSVTFGLDKNDLLFRNLKMTKSSMNPIMAAQCFCDSLS-- 574
Db 572 GKQBAHPSLTHYAGTVYDYNILGWLEKNKOPNETVVALYQKSUKMATLFSVATADTG 631
QY 575 -----KKRP---ETVATQFMSLLOLVEILRSKPEPAVIRCIKENDAKQGRFDEVL 623
Db 632 DSGSKGKGGKGSFQVSAHRENKMLNTNLTTHPHFVRCIIPNERKAPGMDNPLV 691
QY 624 RHQVKYLGMLNLRVRAGFAYRRKYAEFLQRYKSLCPETWPMWAGR---PQDGVAVLVR 680
Db 692 MHQLRCNGVLEGIRICRKGFPNRLYDQFRQRYILNPVLP--EGQFIDSRKGTCKLS 749
QY 681 HLGVKPEYKMGRTKIFIRPKTLFATEDSLVRROSLATKIQAAMRGFHWKQFLRV-- 738
Db 750 SLIDHNOYKFGHTKVFVK-AGLIGLEEMRDERLSRIITRMOAQAGQLMRTEFKKIVE 808
QY 739 KRGAICIQSW 748
Db 809 RRDALLVIQW 818

RESULT 2

US-08-875-435B-4
; Sequence 4, Application US/08875435B
; Patent No. 6593304
; GENERAL INFORMATION:
; APPLICANT: Hasegawa, Kazuhide
; APPLICANT: Arakawa, Emi
; APPLICANT: Oda, Shoji
; APPLICANT: Matsuda, Yururu
; APPLICANT: Takahashi, Katsuhito
; APPLICANT: Sugahara, Michihito
; APPLICANT: Ishiyama, Haruo
; TITLE OF INVENTION: RECOMBINANT DNA COMPRISING DNA CODING
; TITLE OF INVENTION: FOR MYOSIN HEAVY CHAIN SM1 ISOFORM PROTEIN INSERTED INTO
; TITLE OF INVENTION: VECTOR DNA, MICROORGANISM CARRYING THE RECOMBINANT DNA, AND
; TITLE OF INVENTION: AN AGENT FOR TREATMENT OF ARTERIOSCLEROSIS COMPRISING THE
; TITLE OF INVENTION: RECOMBINANT DNA
; FILE REFERENCE: 07898-013001
; CURRENT APPLICATION NUMBER: US/08/875, 435B
; CURRENT FILING DATE: 1997-07-25
; PRIOR APPLICATION NUMBER: PCT/JP96/00134
; PRIOR FILING DATE: 1996-01-25
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 1972
; TYPE: PRT
; ORGANISM: Oryctolagus cuniculus
US-08-875-435B-4

Query Match 22.0%; Score 1190; DB 4; Length 1972;
Best Local Similarity 29.6%; Pred. No. 1-2e-103;
Matches 317; Conservative 196; Mismatches 357; Indels 200; Gaps 30;

QY 40 SEAAFTENLRPRRENLIYTIYIGVLVSNPYDLOIYRSQRMERYGVSVFVPPHFLFA 99
Db 97 NEASVLNLRERYFSGLIYTSGLFCVWNPYKQLPIYSEKIVDMYKKGKHEMPHIVA 156
QY 100 VADTVRALTRERDQAVMI SGESGAGKTEATKRLQFYAETCPAERG-----GAVR 152
Db 157 IADATYRMLQDREDQSILCTGESGAGKTENTKKVIOYLA-VVASHHKGGKDTSTGELE 215

QY 153 DRLLQSNPULFARCNAKTIRLNDSSRFKYMVOVDFKGPVGGHILSYLLEKSRVVHON 212
Db 216 KQLLOANPILFARCNAKTIRLNDSSRFKIRINFDVTGYIVGANIETYLLEKSRIRQA 275
QY 213 HGERNEHFVQQLLEGGBEETLRLGLERNPQSVLYLVKGQCAKVSSINDKSDKWKVKAL 272
Db 276 REERTTHIFYLLAGAKKRWNDLLLE-GFNNTYFLSNG-FVPIPAQDDMEQETVEAM 333
QY 273 SVTDFTDEVEDLLSIVASVLHIGNIHPAEDSN-AQVTTENQLKYLTRLGLGVEGFTL- 330
Db 334 SIMGFSEEEQLSVLVKVVSVLQNLGNIVFKKERTDQASMPDNTAAQKVCHLMGINVTDT 393
QY 331 REALTHKIIAKEGEILLSPNL-EQAYARDALAKAVYSRTFTWLVRKINRSIASKDAES 389
Db 394 RSLTTPR--IKVGRDVVQKATQKEQADFVAALAKATYERLFEWILSRVKNKALDKTHROG 451
QY 390 PSWRSTTVLGLDIYGFVFOHNSPFOFCINYNKELQQLFIELTKSQEYEEAEGIAW 449
Db 452 ASF-----LGILDIAGFEIPEVNSFEQLCINTNEKLOQLFNHTMFIHQEYQREGIEW 506
QY 450 EPVQYNNKIIC-DLVEEKFK-GIISILDECLRGEATDITLFEKLEDVTKPHPHFTL 506
Db 507 NFIDGLDLQPCIELIETERNPNPPGVALLDEBCWPP-KATDKSFEVKLCTEQGNHPKF-- 563
QY 507 HKLADQKTRKSLDRGFRLLHYAGEVYTSVTFGLDKNDLLFRNLKMTKSSMNPIMAAQC 566
Db 564 -----QKPKLKDKTFFSIHVGKVDYNASAWLTKNDPLNDNVTSILNASDSKFEVADL 618
QY 567 F-----DKSELSDKKRP-----ETVATQFMSLLOLVEILRSKPEPAVIR 606
Db 619 WKDVDRIVLGDMQAKMTESLPSASKTKKGMFTVQGLYKEQGLKMLTIRNTTNPVRC 678
QY 607 IKPDNAKQGRDEVLIRHQVKYLGMLNLRVRAGFAYRRKYAEFLQRYKSLCPETWPM 666
Db 679 IIPNHEKRSKGLDAFLVLEQLRCNGVLEGIRICRQGFNRIVQEPFRQRYEILANAIPK 738
QY 667 MAGRPQDGVAVLVRHLGYKPEEYKMGRTKIFIRPKTLFATEDSLVRROSLATKIQAAM 726
Db 739 GEMDGKQACILMIKALELDPNLYRIGQSKIFPR-TGVLAHLEBERDLKITDVIMAFQAMC 797
QY 727 RGFHWKQKFLRVKRSACIQ-----SWWR----- 750
Db 798 RGYLARKAKAKQKQQLTAMKVLRNCAAVLKLNRNQMWFLLTKVXPELLQVTRQEEEMQAK 857
QY 751 -GTLGRKKAARKAAQTIRRIIRGFIIRHSRCPENAEFLDHVRASFLNLRRLPRNV 809
Db 858 EDELQKIKERQOK-ABSEIQLQO-----KHTQLSBE-----KML 891
QY 810 LDTSWPTPPALREASELLRELCKMNMVWKYCRSISPEWKQOLQOKAVASEIFKGGKDN 869
Db 892 LQEQLOAETELYAEABEMVRLAAK-----KQLEEE--ILHE----- 926
QY 870 POSVPRFLFISTRIGTEISPRVLQSLGSEPIQYAVVVKYDRKGYKPRQLLTPSAVV 929
Db 927 -----MEARLEBEDRGQQLQA-----ERK--KMAQQLDLEEQ--- 958
QY 930 IVEDAKVKQRIYANLTG-ISVSSLSLSLFLVHVRQEDNKQKGDVVLQSDHVIETLTATA 988
Db 959 LEEEAARQKLEKVTAEAKIKLEDDILVMDQ--NNKLSKERKLEERISDLTTLNLA 1016
QY 989 LSADRVNNINQSGSITFAGGPRGDIIDFTSGSELLITKAKNGHAAVA 1038
Db 1017 EEEKAKN-----LYKKNKSHSMIS 1037

RESULT 3

US-08-875-435B-3
; Sequence 3, Application US/08875435B
; Patent No. 6593304
; GENERAL INFORMATION:
; APPLICANT: Hasegawa, Kazuhide
; APPLICANT: Arakawa, Emi
; APPLICANT: Oda, Shoji

```

; APPLICANT: Matsuda, Yuzuru
; APPLICANT: Takahashi, Katsuhito
; APPLICANT: Sugahara, Michihiro
; APPLICANT: Ishiyama, Haruo
; TITLE OF INVENTION: RECOMBINANT DNA COMPRISING DNA CODING
; TITLE OF INVENTION: FOR MYOSIN HEAVY CHAIN SM1 ISOFORM PROTEIN INSERTED INTO
; TITLE OF INVENTION: VECTOR DNA, MICROORGANISM CARRYING THE RECOMBINANT DNA, AND
; TITLE OF INVENTION: AN AGENT FOR TREATMENT OF ARTERIOSCLEROSIS COMPRISING THE
; TITLE OF INVENTION: RECOMBINANT DNA
; FILE REFERENCE: 07898-013001
; CURRENT APPLICATION NUMBER: US/08/875,435B
; CURRENT FILING DATE: 1997-07-25
; PRIOR APPLICATION NUMBER: PCT/JP96/00134
; PRIOR FILING DATE: 1996-01-25
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 3
; LENGTH: 1972
; TYPE: PRT
; ORGANISM: Mus musculus
US-08-875-435B-3

```

Query Match	21.7%	Score 1169.5;	DB 4;	Length 1972;
Best Local Similarity	29.8%	Pred. No. 1.1e-101;		
Matches 306;	Conservative	196;	Mismatches 359;	Indels 165; Gaps 26;

QY	40	SEAAFTENURRPRENLIIYITG	PVLVSNVPYRDQIYSRQHMERYAGSVFVPPHLFA	99	
Db	97	NEASVLNHLRERYFSGLIYTY	SGLFVWVNPVKYLPYSEKIVDMYKGRHEMPPIYA	156	
QY	100	VADTVVRLRTRRRDQAVMIS	GESGAGKTEATKRLLOFYAETCAPERG	152	
Db	157	IADTAYRMLQOREDSILCTES	GAGTENTQKVIQYLA-VWASSHKGKXDSITGELE	215	
QY	153	DRLLQSNPVLEAFGNAKTLR	NDNSRFGYKMDVDFDKGAPVGGHILSYLLEKSRVVHQ	212	
Db	216	KOLLQANPILEAFGNAKTV	KNDNSRFGKFRINFDTVIGYVIGANIETYLEKSRAIRQA	275	
QY	213	HOERNHFVYQLLEGEBETLR	RLGLERNPOSYLYLVKQC	KAVSSINDKSDMKWRKAL	272
Db	276	RDERTFHPYLLAGAKMKSD	LILLE-SFNSYTFLSNG-FVP	IPAAQDDMEFQETLEAM	333
QY	273	SVIDTEDEVEDLLISAVSL	HLGNIHFAADEDNS-AQVT	TENOLKYLTRLGLVGGTTLR	331
Db	334	SIMGNESEQUALIKVWSVL	QLGNIYFKKERNTDQASMP	DNTAAQVCHLVGINVTDTF	393
QY	332	EALTRHKIIAKGEEILLS	PLNLEQAAAYARDALAKAV	YSRTFTWLVRKLNRSILASKDAESPS	391
Db	394	RAILTRIKVGEDVVQAKAT	KEQADFALAKAKTYERLFR	WILSRNKALDKTHRQGAS	453
QY	392	WSTTVLGLDIOYGEFVFOH	NSFEQFCINYCNEKLQOLF	IELTLKSPQEEYAEAGTAWEP	451
Db	454	F-----LGILDIAGEFI	FEVNSFEQLCINYTNEKL	QQLFNHTWFLBQEEYQREGIEWNF	508
QY	452	VQY-FNNKIIDLVKEKFK--	GIISIDEECLRGEATD	TFLEKBDTVKPHFLTHK	508
Db	509	IDFGDLQFSHELIERPNN	PFGVIALIDEECWFP-KAT	DKSFVEKLCSEQGNPKF----	563
QY	509	LADQTRKSLDRGEPRLAHY	AGEVTVSYTGFLDKNDLL	FRNLKETWCSSWNFIMAQCF-	567
Db	564	---QPKQLKDKTEFSII	HYAGKVDYNASAWLTON	MDPLNDNVTSLNASDRFVADLMK	620
QY	568	-----DKSELSDKKRP-	-----ETVATQFKWSLL	QLQVLVEILSRKEPAYIRCIK	608
Db	621	DYDRIVGLDQAKMTES	SLPSASKTKGMFTVQQLY	KEQGLKMLATLRNTTANFVRCII	680
QY	609	PNDAKQPGRDEVILRHQVY	KLGLMENLVRRRAGFAYR	KRYEAFQRYKSLCPTWPMWA	668
Db	681	PNHEKRSKGLDAFLVLE	QLRCNGVLEGIRICROG	FPNRIYVQFQRQRYEILANALPKGF	740
QY	669	GEPOGVAIVRHLYKPEEY	KMGRTKIFRFPKTLFAT	EDSILEVRRQSLATKIQAAWRG	728
Db	741	MDGQACATIMIKALELD	PNLYRIQOSKIFFR-TGL	VAHLEERDLKTTVDVMAFOAMCRG	799


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Db 1036 RQFHLHQAQSVIIQRFWRNYLNQKQVDAADAVQKDAFYMAAALLOASWRAHLERQYL 1095
QY 737 RVKRSALICIQSWRGTLKRRKAA---KRKAA---QTIRRLI-----RGFTLR 778
Db 1096 ELRAAAIVIQKWRDYRRHMAAICIQARKKAVRESKRYOEQRKKIILLQSTCRGFRAR 1155
QY 779 HSPRC-----PENAFFLDHVRASFLNLRRQLPRNVLDTSWPTPPPALREASBELL 828
Db 1156 QRKALKAEQRLRETKPEVG--LVNIKGYSLIQSGSDPSEMEDCSFDNRKAIKAECKSVI 1213
QY 829 RELCKMVMVYCRSISPEWKOQLOQKAVASEIPKGGKDNYPQSVPRLFSTRLGTBEIS 888
Db 1214 ESNRISRESSVDCLKESPNKQOERASQSGVDL---QED-----VLVRER----- 1255
QY 889 PRVLQSLGSEPIQYAVPVVVKYDRKGYKPRPQLLLTSAVVIVEDAKVKQRIDYANLTGI 948
Db 1256 PRSLEDLHQQKVGRA-----KRESRMELEQAIFSLLLKVR-----SLGGI 1298
QY 949 SVSLSLSLFLVHQREDNKQGDVW---LQSDH-----VIETLTKTALSADRVN 995
Db 1299 SPS-----EDRRWSTELVPEGLQSPRGTPDSESSQGSLELLSYEEBSQKSKLE 1345
QY 996 NININOISITF 1006
Db 1346 SVISDEGDLQF 1356

RESULT 7
US-09-883-134-4
; Sequence 4, Application US/09883134
; Patent No. 6511840
; GENERAL INFORMATION:
; APPLICANT: Walke, D. Wade
; APPLICANT: Scoville, John
; APPLICANT: Donoho, Gregory
; APPLICANT: Turner, C. Alexander Jr.
; APPLICANT: Mathur, Brian
; APPLICANT: Mathur, Daniel
; APPLICANT: Friddle, Carl Johan
; TITLE OF INVENTION: No. 6511840el Human Kinase Proteins and Polynucleotides Encoding
; FILE REFERENCE: LEX-0193-USA
; CURRENT APPLICATION NUMBER: US/09/883,134
; CURRENT FILING DATE: 2001-06-15
; PRIOR APPLICATION NUMBER: US 60/211,572
; PRIOR FILING DATE: 2000-06-15
; PRIOR APPLICATION NUMBER: US 60/216,382
; PRIOR FILING DATE: 2000-07-07
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 1236
; TYPE: PRT
; ORGANISM: homo sapiens
US-09-883-134-4

Query Match 20.08; Score 1078.5; DB 4; Length 1236;
Best Local Similarity 33.08; Pred. No. 2.5e-93;
Matches 260; Conservative 161; Mismatches 284; Indels 83; Gaps 17;

QY 31 DFVLENTSEAAFIENLRFRFRNLIYTVIGPVLVSVNVPYRDLQIYSRQHMERYGVSP 90
Db 335 DLVNLV-FLVDEDTIHQKRYADLLIYTVGDLIALNFPQNLISYSPQFSRLYHGVR 393
QY 91 YEVPPLHFAVADTVYRALRTRRDQAVMISGESGAGKTEATKLLQFYAETCPAPERGGA 150
Db 394 ASNPPIHIFASADAAYQCMVTLTKDQCVISGESGSGKTESAHLIVQHL--TELKANNQT 451
QY 151 VRDLQSNPVLFAFGNAKILRNDNSRFGKYMVDQDFKAPGVGHILSYLLEKRVVH 210
Db 452 LREKILQVNLFAFGNSCTAINDNSSRFKYLEMFTPTGVVMGARISYLLLEKSRVIK 511
QY 211 QNHGERNFHYFQYLLLEG-GEETLRLUGL-ERNPQSVYLKVGQCAKVVSSINDKSNKVM 268

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Db 512 QAAREKNFHIFYIYAGLHHQKLSDFRLPEEKPPRYADTGEV--MHDITSKESYRRQ 569
QY 269 RKALS-----VIDFTEDVEDLLSTIVASVLGNHFAA-----DEDSNAQVTTENQLKYL 320
Db 570 FEAIQHCFRIIGETDKVHSVYRILAGILNIGNIEFAAISQHOTDKSEVPNAEALQNA 629
QY 321 RLIGVEGTTUREALTTHRKIIAKGBELLSPNLQAAVARDALAKAVYSRTFTMLVRKINR 380
Db 630 SVLCISPEEQEALTSCHVVRGETTIIRANTVRADVRDAMSKALGRUFSWIVNRIN- 688
QY 381 SLASKOAESPSWSTTVLGLLDIYGFEVFOHNSFEQFCINYCNKELQQLFIETLTKSE 440
Db 689 TLLQDENICAGSGMNVGILDFGFENFQNSFEQLCINIANEQIQYFNFQHVFALEQM 748
QY 441 EYAEAGTIAMEPVOYFNKIIICDLVEKFKGIISILDEECLRPGBATDLTPEKLEDTVKP 500
Db 749 EYQNEGIDAIPVEYEDNRPLDMFLQPLGLLALLDEESRFP-QATDQTLVDKFDNLR- 806
QY 501 HPHFLTHKLADQKTRKSLDRGEPRLLHYAGEVYVTVGFELDKNDLLFRNLKETMCSMN 560
Db 807 CKYFWRPKGVELC-----FGIQHYACKVLYDASGVLEKRDITLPADVVVVLRITSE 857
QY 561 PIMAQCFD-----KSELSDKKRP 578
Db 858 KLLQQLFSIPLTKTGNLAOTRARITVASSSLPPHFSAGAKAVDTLEVIRHPEETNMKR- 916
QY 579 ETVATOPKMSLLQLVELRSKEPAYIRCIKPNDAKQBPGRDEVLIRHGVKYLGMENLRV 638
Db 917 QTVASYFRYSLMDLLSKMVGQPHFVRCIKPNDREALOFSRERVLQAQRSTGILETVSI 976
QY 639 RRAGFAYRRKYEAFLQRYKSLCPETWPMWAGRPQDGVAVLVRHLGYKPEYKMGRTKIFI 698
Db 977 RRQGYSHRILFEFVRYVYLAFTAHTQTPASKESCVAILLEKS---RLDHWVLGKTQVFL 1033
QY 699 RPFKTLFATEDSLEVRQSL--ATKIQAAWRGHFWKQFLRVK---RSAICIQSWWRGT 752
Db 1034 KY----YHVEQLNLLREVIQVGVVVLQAYTKGVLGARYKVKREKREKGAIAIQSAWRGY 1089
QY 753 LGRRKAAK 760
Db 1090 DARRKPK 1097

RESULT 8
US-08-938-105-3
; Sequence 3, Application US/08938105
; Patent No. 6353151
; GENERAL INFORMATION:
; APPLICANT: Leinwand, Leslie A.
; APPLICANT: Vikstrom, Karen L.
; TITLE OF INVENTION: TRANSGENIC MODEL FOR HEART FAILURE
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Sheridan Ross P.C.
; STREET: 1700 Lincoln St., Suite 3500
; CITY: Denver
; STATE: CO
; COUNTRY: U.S.A.
; ZIP: 80203
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/938,105
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Crook, Wannel M.
; REGISTRATION NUMBER: 31,071
; REFERENCE/DOCKET NUMBER: 3595-4
; TELECOMMUNICATION INFORMATION:

```

Query Match 18.9%; Score 1018; DB 4; Length 1885;
Best Local Similarity 32.1%; Pred. No. 3.2e-87;
Matches 253; Conservative 136; Mismatches 286; Indels 114; Gaps 17;

Query Match	10.7%;	Score	575.5;	DB	4;	Length	1695;
Best Local Similarity	23.7%;	Pred. No.	5.3e-45;				
Matches	201;	Conservative	159;	Mismatches	325;	Indels	163;
						Gaps	25;
<hr/>							
QY	38	FTSEAFIENLRFRRENLIYTIYIGVLVSNVNPYRDLOIYRQWHERYRGVSFYEVP	PHL	97			
		: : : : : : : : : : : : : : : : : : :					
Db	75	YLNESVLLTRQYCASLHTYAGPSLLVILSTRGAPAVYSEKVKWHMFKGCRREDMA	PHI	134			
		: : : : : : : : : : : : : : : : : : :					
QY	98	PAYADIVYRALPTFRERDOAWMLSGSGAGKTEATKRLQFYAETCTPAPERGA	VRD	157			
		: : : : : : : : : : : : : : : : : : :					
Db	135	YAAQATAYRAMLMSRDQIQIVILGSSGGSKTTSFQHLVQVLTATIAGTSGTK	VFS	194			
		: : : : : : : : : : : : : : : : : : :					
QY	158	SNPVLAFNGNAKTLRNDNSSREFGKYMDFQFKGAPVCGHILSVILLEKSRV	VQH	217			
		: : : : : : : : : : : : : : : : : : :					
Db	195	LSLTLEAFNGSPITMNGSATRESQTLSDLFQAGOVASASIQTMLEKLRVAR	PASE	254			
		: : : : : : : : : : : : : : : : : : :					
QY	218	FHVFYQLLEGGEETLRRIGLERNPQSYLYLVKGCQAKVSSINDK----	SDK	273			
		: : : : : : : : : : : : : : : : : : :					
Db	255	FNVFYLLACGDAITRTELHNLHNAENNVFGI----VPLSKPEEKAAQOFK	LQAA	310			
		: : : : : : : : : : : : : : : : : : :					
QY	274	VIOFTDEVEDLLSIVASVHLHG--NIHFAADEDSNAQVTTENQKYLTRLG----	324				
		: : : : : : : : : : : : : : : : : : :					
Db	311	VLAISPEEQKTCWLLIASIYHLGAAGATKEAAEGRQFARHEWAQKAAVYLL	GCSE	370			
		: : : : : : : : : : : : : : : : : : :					
QY	325	-----VBGTTLREALTHRK-----IIAKGBEILLSPLNLEQAAVARDALAK	AVYS	372			
		: : : : : : : : : : : : : : : : : :					
Db	371	SAIFXHQKGGTLOKSTSFROQPEESGUGBGTU-----SALCEHGWASGL	YSBEL	423			
		: : : : : : : : : : : : : : : : : :					
QY	373	WLVRKINRSLAKDAESPWSRSTTVLGLDLYGFE-----VFQHNSEFOFC	INCY	426			
		: : : : : : : : : : : : : : : : : :					

371 SAIFKHQLKGGTLQRSTSTFRQPEESGLGEGTKL-----SALECLEGVASGLYSELFT 423

373 WLVRKINRSLASKDAESPSPWRSTTVLGLLDIYGFE-----VFQHSFEQFCINVCNEKL 426

Db 424 LLSLVNRALKS-----SOHSLCSMMIVDTFGQNPWGGSGARGASFEELCHNTAQDRL 477
QY 427 QQLFIETLTKSEOEYEAEAGIA-----WEPVOYEN-----NKIICDLAVE-EKEKGIIS 473
Db 478 QRLFHRTLQELERKEDNIELAFDDLEFVADDSVAADVQASHLVRSLAHADEARGLLM 537
QY 474 ILDEECLRGATDITLTFLEKLEDTVKPHPHLTHKLADQTRKSLDRG-----EFLILHYA 529
Db 538 LLEEEALVPG-ATEDALLDRLFSYGPQE-----GDKKGQSPLLRSSKPRHLLGHSH 589
QY 530 GE--VTYSVTGELD--KNDLLFRNLKETWCSMMPIMAOCF-----567
Db 590 GTNWVEYNVAGWLNITKQNPAT--QNPRLQDSQKRIISNLFGRAGSATVLSSGIAGLE 648
QY 568 DKSELSDDK-----RPETVATQFMSLLQLVLILRSKEPAYIRCIKPND 612
Db 649 GGSQALRRATSMRKTTFTTGMAAVKKSLSICIKLQVDALITIKRSMHFFHCFLPVAE 708
QY 613 KQPG-----RFEVLIRHQVKYLGIMENLVRRAFG 643
Db 709 GWPGEPRSSRRVSSSELDLPPGDPCEAGLLQLDVSLLRAQLRGSRLLDAMRVQGY 768
QY 644 AYRRKYEAFLQYKSLCPETWPMAGR-----PQDGVAVLVRHLGYKPEEYKMGRTKIF 697
Db 769 PDMVFPSEPRRFDVLAPHLTK--HGRNVIYVDEKRAVEELLESLDLEKSSCCLGLSRVF 827
QY 698 IRFPKTLFATESLEVRROSLATKIQAAMRGFWKQFLVRKRSAL---CIQS-----747
Db 828 FR-AGTLARLEQRDEQTSHTLTFQAACRGYLAHQHFKKIKIQLAIRCVQKNKNG 886
QY 748 -----WWR 750
Db 887 VKDWPWMK 894

RESULT 10

US-09-866-108A-15754
; Sequence 15754, Application US/09866108A
; Patent No. 6686188
; GENERAL INFORMATION:
; APPLICANT: GU, Yizhong
; APPLICANT: JI, Yonggang
; APPLICANT: PENN, Sharon G.
; APPLICANT: HANZEL, David K.
; APPLICANT: RANK, David R.
; APPLICANT: CHEN, Wensheng
; APPLICANT: SHANNON, Mark
; TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
; FILE REFERENCE: AEOMICA-7
; CURRENT APPLICATION NUMBER: US/09/866,108A
; CURRENT FILING DATE: 2001-05-25
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; Remaining Prior Application data removed - See File Wrapper or PALM.

; NUMBER OF SEQ ID NOS: 15755
; SOFTWARE: Aecomica Sequence Listing Engine
; Patent No. 6686188
; SEQ ID NO 15754
; LENGTH: 1561
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-866-108A-15754

Query Match 9.6%; Score 519.5; DB 4; Length 1561;
Best Local Similarity 23.5%; Pred. No. 1.1e-39;
Matches 190; Conservative 130; Mismatches 319; Indels 171; Gaps 22;

QY 82 MERVGVFYEVPVPHLFAVADTVVTRALRTERDDQAVMISGESGAGKATEATKRLLOFYAET 141
Db 1 MEMFKGCRREDMAHIYAVAQATAYRAMLSRQDQASILLGSSGSKTSCQHLVQYLATI 60
QY 142 CPAPERGAVDRLLQSNFVLEAFGNAKTLNDNSSRFGKYMVDVQDFKGPAPVGHILSY 201
Db 61 AGISGNKVFSEVKWQALYTLLEAFGNSPTIINGNATRFSQLSLDFDQAGQVASASICTM 120
QY 202 LLEKSRVVHQNHGERNHVFVQLLEGGBEEETRLRLGLERNPQSYLYLVKGCQAKVSSIND 261
Db 121 LLEKLRVARRPASEATFNVFYLLACGGDTLRTTELHNLHNAENVVFGIVPLAKPEEKQA 180
QY 262 KSDMKWKRKALSVIDFTEDEVEDLLSIVASVLHLG--NIHFAADEDSNAQVTTENQKYL 319
Db 181 AQQSKLQAAMKVLGISPDQKACWFLAAIYHLGAAGATKEAAEAGKQFARHEWAKA 240
QY 320 TRLLGVEGTTTREALTHRKIIAKGEELLSPNLBQ-----AAYARDALA 363
Db 241 AYLLGCSLEELSSAIFKHQ--HKGDTLQRSTSFQGPESGLGDGTGPKLSALECLEGMA 298
QY 364 KAVYSRTFTLVRKINRSLSKDAESPSRSTTVLGLLDIYGEFVQHN-----SFEQF 417
Db 239 AGLYSELTLLVSLVNRALKS-----SOHSLCSMMIVDTFGQNPWGGSGARGASFEEL 352
QY 418 CINYNEKLOQLFIETLTKSEOEYEAEAGIA-----MEPV-----QVFNKKIICDL 463
Db 353 CHNYTQDRLQRLFHRTFVQELERYKEBNIELAFDDLEPPTDSDVAADVQASHQSLVRS 412
QY 464 VE-EKFGIILDEECLRGATDITLTFLEKLE-----TVKPHPHL 505
Db 413 ARTDEARGLLWLEEEALVPGASED--TLLERLFYSYGPQEGDKKGQSPLLHSSKPHHFL 471
QY 506 THKLADQKTRSLDRGEPFLHLYAGEVTVSYTGFLD--KNDLLFRNLKETWCSMNPIM 563
Db 472 GHS-----HGTNWVEYNVAGWLNITKQNPAT--QNPRLQDSQKII 512
QY 564 AQCF-----DKSELSDDK-----RPETVATQFMSLL 590
Db 513 SNLFLGRAGSATVLSSGIAGLEGGSQLALRRATSMRKTTFTTGMAVAVKKSLSICIQMLQVD 572
QY 591 QLVLEILRSKEPAYIRCIKP-----NDAKQPG--RPEDEV 621
Db 573 ALIDTIKKKHLHFVHCFLPVAEGWAGPRSSRRVSSSELDLPSGDHCEAGLLQLDVP 632
QY 622 LIRHQVKYLGIMENLVRRAGFAYRRKYEAFLQYKSLCPETWPMAGR-----PODGV 675
Db 633 LLRTQLRGSRLLDAMRVYRQGYPDHMFSEPRRFDVLAPHLTK--HGRNVIYVDERAV 691
QY 676 AVLVRHLGYKPEEYKMGRTKIFIRFPKTLFATESLEVRROSLATKIQAAMRGFWKQF 735
Db 692 EELLECLDLEKSSCCMGLSRVFR-AGTLARLEQRDEQTSRNLTLFQAACRGYLAHQH 750
QY 736 LRVKRSAL---CIQS-----WWR 750
Db 751 KKRKIQLAIRCVQKNKNGVKDWPWMK 780

RESULT 11

US-09-866-108A-3
; Sequence 3, Application US/09866108A

[illegible]

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QY 137 FYAETCPAPERGGAVR--DRLLQS-----NPVLEAFGNKATLRN---DNSS--- 177
Db 382 FDAPEAPLPSRKAALVTRLTSLDFTDYASRIIHPVTRTLDQSPELRSTAMDTLSSLV 441
QY 178 -RFGKYMDVQDFKGA PVGGHLSYLEKSRVHQNH-----GERNFHVFY 222
Db 442 FOLGKKYQIFIP-----MVNKLVRHRIHQRYDVLCIRIVKGYTLADEEDPLIY 492
QY 223 Q--LLEGGEBETLRLGLERNPQSYLYLVKQCAKUSSINDKSDWKVMKALSVIDFTED 280
Db 493 QHRMLRSQGDALASGVETGPMKKLH-----VSTINLQKAWGAARR----- 534
QY 281 EVEDLLSIVASVLHLGNIHFAEDSNQAQVTENQLKYLTRLLGVEGTTLREALTHRKII 340
Db 535 -----VSKDDWLEWLRRL----- 547
QY 341 AKGBELSLPLNLEQAAYARDALAKAVYSRTFTWLVRKINRSLASKDAESPSWSTTVLG- 399
Db 548 --SLELL-----KSSSPSLRSCWALAQ 568
QY 400 ----LLDIYGFVFOHNSFEQFCINCEKLOQLF---IELTLKSEQEEYEAEGIAWEP 451
Db 569 AYNPMARDLF-----NAAFVSCWSELNEDQOQDELIRSIELALTSQD-----IA-EV 613
QY 452 VOYFNKKIICDLVEEKFKGIISILDE-----ECLRPGEATDLTFLEKLEDTVKP 500
Db 614 TOTLLN--LAEPMEHSDKGPLPLRDNDGIVILGERAAKCRAYAKA---LHYKELEFQKGP 668
QY 501 HPFEL-----THKLADQKTRKSLDRGEFRLLHYAGVETYSVTGF----- 539
Db 669 TPAILESLSINNNK--QOPEAAAGVLEYAMKHF-GELEIQATWYEKLHEWEDALVAYDK 725
QY 540 -LDKNNDLFRNLKETMCSMNPIA-----OCFDKSELSDKKRPETVATQFMKSL 589
Db 726 KMDTNKD--DPELMGLGRMCLEALGEGWQLHQCCERKWTLVN---DETQAKMARMAA 777
QY 590 LQVLVELIRSKEPAYIRICIKNDKQKPRFDEVLIRHQ-----VKYLGIMENLRVR 639
Db 778 AAANGLGQWDSMEEYTCMIPRTHDGA FYRAVLALHQDLFSLAQOQCIDKARDLLDAELTA 837
QY 640 RAGFYARKYAEFLQRYKSCPETWPMWAGPQDGVAVLVRHLGYKPEE---YKMGRTKI 696
Db 838 MAGESYRAYGAMVS-----C-----HMLSELEEVIOYKL----- 867
QY 697 FIRPKTLFATEDSLEVRROSLATKIQAAWGFHWRQKFLVRKRSALC-----IQSWRGT 752
Db 868 -----VPERREIIRQIWWERLQCCQRIVEDQKILMVRSLVSPHEDMRTW----- 913
QY 753 LGRKKAARKWAAQTIRRLTRGFI LRHSRPCPENA--FFLDHVRASFLNLRRLPRNVLD 811
Db 914 -----LKYASLCGSGRLAHLAKTILVLLGVD---PSRQLD 946
QY 812 TSWPTPPALREASELLRCLMKNVMYKCRSISP-----EMKQOOLQOKAVASEIFK 863
Db 947 HPILPTVHPQTYA-----YKQNM--WKSARKIDAFQHMQHVFQTMQOQAHAIAETDQ 998
QY 864 GKNDYQSVPRPFI STRLGTEBISPRVLOSLGSEPI-----QYAPVVKYDRGKYPKPR 919
Db 999 HKQELH-KLMARCFI--KLGEWQLN--LOGINESTIPKVLQYYSAAETHDRSRYKAWHA 1052
QY 920 QLLLTTSANV-----IVEDAKYQOR-IDYANITGHSVLSLSLFLVHLVQREDNKQGD 972
Db 1053 WAYMNFPAVLHYKHQONQARDEKKLRHAGSANTTNATTAATTAATTASTEGSSESE 1112
QY 973 VLIQSD-----HVIEFTLTKT-----ALSADRNNI 997
Db 1113 AESTENSPSPLOKKVTEDLSTLLMYTPVAVQGFERSISLRGNL 1160
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RESULT 14

US-09-012-504A-12

; Sequence 12, Application US/09012504A

; Patent No. 6464974

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; GENERAL INFORMATION:
; APPLICANT: Berlin, V.
; APPLICANT: Chiu, I.
; APPLICANT: Cottarel, G.
; APPLICANT: Damagnez, V.
; TITLE OF INVENTION: IMMUNOSUPPRESSANT TARGET PROTEINS
; FILE REFERENCE: APBI-P05-036
; CURRENT APPLICATION NUMBER: US/09/012,504A
; CURRENT FILING DATE: 1998-01-23
; PRIOR APPLICATION NUMBER: 08/360,144
; PRIOR FILING DATE: 1994-12-20
; PRIOR APPLICATION NUMBER: 08/250,795
; PRIOR FILING DATE: 1994-05-27
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: Patent in version 3.1
; SEQ ID NO 12
; LENGTH: 1809
; TYPE: PRT
; ORGANISM: Mammalian
; US-09-012-504A-12
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Query Match 3.0%; Score 160; DB 4; Length 1809;
Best Local Similarity 19.1%; Pred. No. 3.2e-05;
Matches 193; Conservative 121; Mismatches 318; Indels 376; Gaps 46;

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QY 137 FYAETCPAPERGGAVR--DRLLQS-----NPVLEAFGNKATLRN---DNSS--- 177
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QY 178 -RFGKYMDVQDFKGA PVGGHLSYLEKSRVHQNH-----GERNFHVFY 222
Db 442 FOLGKKYQIFIP-----MVNKLVRHRIHQRYDVLCIRIVKGYTLADEEDPLIY 492
QY 223 Q--LLEGGEBETLRLGLERNPQSYLYLVKQCAKUSSINDKSDWKVMKALSVIDFTED 280
Db 493 QHRMLRSQGDALASGVETGPMKKLH-----VSTINLQKAWGAARR----- 534
QY 281 EVEDLLSIVASVLHLGNIHFAEDSNQAQVTENQLKYLTRLLGVEGTTLREALTHRKII 340
Db 535 -----VSKDDWLEWLRRL----- 547
QY 341 AKGBELSLPLNLEQAAYARDALAKAVYSRTFTWLVRKINRSLASKDAESPSWSTTVLG- 399
Db 548 --SLELL-----KSSSPSLRSCWALAQ 568
QY 400 ----LLDIYGFVFOHNSFEQFCINCEKLOQLF---IELTLKSEQEEYEAEGIAWEP 451
Db 569 AYNPMARDLF-----NAAFVSCWSELNEDQOQDELIRSIELALTSQD-----IA-EV 613
QY 452 VOYFNKKIICDLVEEKFKGIISILDE-----ECLRPGEATDLTFLEKLEDTVKP 500
Db 614 TOTLLN--LAEPMEHSDKGPLPLRDNDGIVILGERAAKCRAYAKA---LHYKELEFQKGP 668
QY 501 HPFEL-----THKLADQKTRKSLDRGEFRLLHYAGVETYSVTGF----- 539
Db 669 TPAILESLSINNNK--QOPEAAAGVLEYAMKHF-GELEIQATWYEKLHEWEDALVAYDK 725
QY 540 -LDKNNDLFRNLKETMCSMNPIA-----OCFDKSELSDKKRPETVATQFMKSL 589
Db 726 KMDTNKD--DPELMGLGRMCLEALGEGWQLHQCCERKWTLVN---DETQAKMARMAA 777
QY 590 LQVLVELIRSKEPAYIRICIKNDKQKPRFDEVLIRHQ-----VKYLGIMENLRVR 639
Db 778 AAANGLGQWDSMEEYTCMIPRTHDGA FYRAVLALHQDLFSLAQOQCIDKARDLLDAELTA 837
QY 640 RAGFYARKYAEFLQRYKSCPETWPMWAGPQDGVAVLVRHLGYKPEE---YKMGRTKI 696
Db 838 MAGESYRAYGAMVS-----C-----HMLSELEEVIOYKL----- 867
QY 697 FIRPKTLFATEDSLEVRROSLATKIQAAWGFHWRQKFLVRKRSALC-----IQSWRGT 752
Db 868 -----VPERREIIRQIWWERLQCCQRIVEDQKILMVRSLVSPHEDMRTW----- 913
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Qy 753 LGRKAAKRWAAQTIRRLIRGFILRHSPCPENA-FFLDHVRASFLNLRRLQPRNVL 811
Db 914 -----LKYASLCKGSRGLAHLKTLVLLGLVD---PSRLD 946
Qy 812 TSWPTPPALREASELLRELCKMVMWKYCRSISP-----EWKQLOQKAVASEIFK 863
Db 947 HPLPTVHPQVTYA-----YMKQM-WKSARKIDAFQHMHFVQTMQOQAHAIAITEDQ 998
Qy 864 GKDNYPQSVPRFLFSTRLGTBEISPRVLOSLGSEPI---QYAVPVVVKYDRKGKPRR 919
Db 999 HKQELH-KLMARCF--KLGEWOLN---LQGINESTIPKVLQYISAATEHRSYKAWHA 1052
Qy 920 QLLLTPSAVV-----IVEDAKVKQR-IDYANLTGISVSLSDSLFVLHVQREDNKOKGD 972
Db 1053 WAVNFEAVLHYKHQNAQDEKKLKHASGANITNATTAATTAATTASTEGSSESE 1112
Qy 973 VVLQSD-----HVIETLTKT-----ALSADRVNMI 997
Db 1113 AESTENSPTPSPLOKKVTEDLSKTLMTYVPAVQGFPRFSISLSRGNL 1160

RESULT 15
US-09-012-399A-12
; Sequence 12, Application US/09012399A
; Patent No. 6509152
; GENERAL INFORMATION:
; APPLICANT: Berlin, Vivian
; APPLICANT: Chiu, Maria Isabel
; APPLICANT: Cottarel, Guillaume
; APPLICANT: Damagnez, Veronique
; TITLE OF INVENTION: IMMUNOSUPPRESSANT TARGET PROTEINS
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FOLEY, HOAG & ELIOT LLP
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02109-2170
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/012.399A
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/360,144
; FILING DATE: 20-DEC-1994
; ATTORNEY/AGENT INFORMATION:
; NAME: Vincent, Matthew P.
; REGISTRATION NUMBER: 36,709
; REFERENCE/DOCKET NUMBER: APV-036.02
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-832-1000
; TELEFAX: 617-832-7000
; INFORMATION FOR SEQ ID NO: 12:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1809 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-09-012-399A-12

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Query Match 3.0%; Score 160; DB 4; Length 1809;
Best Local Similarity 19.1%; Pred. No. 3.2e-05;
Matches 193; Conservative 121; Mismatches 318; Indels 376; Gaps 46;

Qy 137 FYAETCPAPERGGAVR--DRLLQS-----NPVLEAFGNKTLRN---DNSS--- 177
Db 382 FDAEALPLSKAALETVDRLTESLDTDYASRIIHFIVRTLQDSPELRSTAMDTLSLV 441

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Qy 178 -REGKYMVDQDFKGPVPGGHIIISYLLEKSRVVQH--GERNFVFY 222
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Qy 223 Q--LLEGGESETLRLGLERNPOSILYLVKQGCAKVVSSINDKGDWKVMRKALSVIDFTED 280
Db 493 QHRMLRSQGDALASGPVETGPMKKLH-----VSTINLQKAWGAARR----- 534
Qy 281 EVEDLLISIVASVLHLGNIHFAADEDSNAQVTTNQLKYLTRLGLGVEGTTLREALTHRKII 340
Db 535 -----VSKDDWLEWLRL----- 547
Qy 341 AKGEBLLSPNLQAAVARDALAKAVYSRTFTVLVRKINRSLASKDAESPSWSTTVLG- 399
Db 548 --SLELL-----KSSSPSLKSCWALAQ 568
Qy 400 -----LLDIYGFVFOHNSFEQFCINYCNKLOQLF---IELTKSEQEEYEAGIAWEP 451
Db 569 AYNPMARDLF-----NAAFVSCWSELNEDQDELIRSLIELALTSDQ-----IA-EV 613
Qy 452 VOYFNKKIICDLVBEKPKGIISILDE-----ECLRPGBATDLTLEKLEDTVKP 500
Db 614 TOTLLN--LAEFMEHSDKGPLRLDDNGIVLLGERAAKCRAYAKA---LHYKSELEFQKGP 668
Qy 501 HPHFL-----THKLADQKTRKSLDRGEPRLLHVAGEVTVSTGF----- 539
Db 669 TPAILESLSINNKL--QOPEAAAGVLEYAMKHF-GELEIQATWYKLEHWEALVAYDK 725
Qy 540 -LDKNNDLLFRNLKETWCSSMNPIMA-----QCFDKSELSDKRPETVATQFMSL 589
Db 726 KMDTKD-----DPFLMLGRMRCLEALGEWQLHQCCCKEKTLVN---DETQAKMARMAA 777
Qy 590 LQVLEILRSKEPAYIRCIKPNDAKQGRDPEVLIRHQ-----VKVLGLMENLRVR 639
Db 778 AAAGLQWDSMEEYTCMTPRDTHDGAFYRAVLALHQLDLSLAQQCIDKARDLLDELTA 837
Qy 640 RAGFAYRRKYEAFLQRYKSLCPETPMWAGRPDGVAVLVRHLGYKPEE---YKMGRTKI 696
Db 838 MAGESYSRAYGAMVS-----C-----HMLSELEEVIOYKL----- 867
Qy 697 FIRFPKTLFATEDSLVRRQSLATKIQAARWGFHWRQKFLRVKRSATC-----TQSWWRGT 752
Db 868 -----VPERREIIRQIWWERLQGCQRIVEDQWKILMVRSLVSVSPHEDMRTW--- 913
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Qy 812 TSWPTPPALREASELLRELCKMVMWKYCRSISP-----EWKQLOQKAVASEIFK 863
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Qy 864 GKDNYPQSVPRFLFSTRLGTBEISPRVLOSLGSEPI---QYAVPVVVKYDRKGKPRR 919
Db 999 HKQELH-KLMARCF--KLGEWOLN---LQGINESTIPKVLQYISAATEHRSYKAWHA 1052
Qy 920 QLLLTPSAVV-----IVEDAKVKQR-IDYANLTGISVSLSDSLFVLHVQREDNKOKGD 972
Db 1053 WAVNFEAVLHYKHQNAQDEKKLKHASGANITNATTAATTAATTASTEGSSESE 1112
Qy 973 VVLQSD-----HVIETLTKT-----ALSADRVNMI 997
Db 1113 AESTENSPTPSPLOKKVTEDLSKTLMTYVPAVQGFPRFSISLSRGNL 1160

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Search completed: March 30, 2004, 15:13:36
Job time : 41.4868 secs